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(11) EP 1 074 617 A2

(12)

EUROPEAN PATENT APPLICATION

(43) Date of publication: 07.02.2001 Bulletin 2001/06

(21) Application number: 00116126.4

(22) Date of filing: 28.07.2000

(51) Int Cl.7: **C12N 15/12**, C12N 15/11, C07K 14/47, C07K 16/18, C12Q 1/68

(84) Designated Contracting States:

AT BE CH CY DE DK ES FI FR GB GR IE IT LI LU MC NL PT SE

Designated Extension States: AL LT LV MK RO SI

(30) Priority: 29.07.1999 JP 24803699

27.08.1999 JP 30025399 11.01.2000 JP 2000118776 02.05.2000 JP 2000183767 09.06.2000 JP 2000241899

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(54) Primers for synthesising full-length cDNA and their use

(57) Primers for synthesizing full-length cDNAs and their use are provided.

5602 cDNA encoding a human protein has been isolated and nucleotide sequences of 5'-, and 3' -ends of the cDNA have been determined. Furthermore, prim-

ers for synthesizing the full-length cDNA have been provided to clarify the function of the protein encoded by the cDNA. The full-length cDNA of the present invention containing the translation start site provides information useful for analyzing the functions of the protein.

Description

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FIELD OF THE INVENTION

5 [0001] The present invention relates to a polynucleotide encoding a novel protein, a protein encoded by the polynucleotide, and new uses of these.

BACKGROUND OF THE INVENTION

[0002] Currently, the sequencing projects, the determination and analysis of the genomic DNA of various living organisms have been in progress all over the world. The whole genomic sequences of more than 10 species of prokaryotes, a lower eukaryote, yeast, and a multicellular eukaryote, C. elegans are already determined. As to human genome, which is supposed to be composed of three thousand million base pairs, the world wide cooperative projects have been under way to analyze it, and the whole structure is predicted to be determined by the years 2002-2003. The aim of the determination of genomic sequence is to reveal the functions of all genes and their regulation and to understand living organisms as a network of interactions between genes, proteins, cells or individuals through deducing the information in a genome, which is a blueprint of the highly complicated living organisms. To understand living organisms by utilizing the genomic information from various species is not only important as an academie subject, but also socially significant from the viewpoint of industrial application.

[0003] However, determination of genomic sequences itself cannot identify the functions of all genes. For example, as for yeast, only the function of approximately half of the 6000 genes, which is predicted based on the genomic sequence, was able to be deduced. As for human, the number of the genes is predicted to be approximately one hundred thousand. Therefore, it is desirable to establish "a high throughput analysis system of the gene functions" which allows us to identify rapidly and efficiently the functions of vast amounts of the genes obtained by the genomic sequencing.

[0004] Many genes in the eukaryotic genome are split by introns into multiple exons. Thus, it is difficult to predict correctly the structure of encoded protein solely based on genomic information. In contrast, cDNA, which is produced from mRNA that lacks introns, encodes a protein as a single continuous amino acid sequence and allows us to identify the primary structure of the protein easily. In human cDNA research, to date, more than one million ESTs (Expression Sequence Tags) are publicly available, and the ESTs presumably cover not less than 80% of all human genes.

[0005] The information of ESTs is utilized for analyzing the structure of human genome, or for predicting the exonregions of genomic sequences or their expression profile. However, many human ESTs have been derived from proximal regions to the 3'-end of cDNA, and information around the 5'-end of mRNA is extremely little. Among these human cDNAs, the number of the corresponding mRNAs whose encoding protein sequences are deduced is approximately 7000, and further, the number of full-length therein is only 5500. Thus, even including cDNA registered as EST, the percentage of human cDNA obtained so far is estimated to be 10-15% of all the genes.

[0006] It is possible to identify the transcription start site of mRNA on the genomic sequence based on the 5'-end sequence of a full-length cDNA, and to analyze factors involved in the stability of mRNA that is contained in the cDNA, or in its regulation of expression at the translation stage. Also, since a full-length cDNA contains ATG, the translation start site, in the 5'-region, it can be translated into a protein in a correct frame. Therefore, it is possible to produce a large amount of the protein encoded by the cDNA or to analyze biological activity of the expressed protein by utilizing an appropriate expression system. Thus, analysis of a full-length cDNA provides valuable information which complements the information from genome sequencing. Also, full-length cDNA clones that can be expressed are extremely valuable in empirical analysis of gene function and in industrial application.

[0007] Therefore, if a novel human full-length cDNA is isolated, it can be used for developing medicines for diseases in which the gene is involved. The protein encoded by the gene can be used as a drug by itself. Thus, it has great significance to obtain a full-length cDNA encoding a novel human protein.

[0008] In particular, human secretory proteins or membrane proteins would be useful by itself as a medicine like tissue plasminogen activator (TPA), or as a target of medicines like membrane receptors. In addition, genes for signal transduction-associated proteins (protein kinases, etc.), glycoprotein-associated proteins, transcription-associated proteins, etc. are genes whose relationships to human diseases have been elucidated. Moreover, genes for disease-associated proteins form a gene group rich in genes whose relationships to human diseases have been elucidated.

[0009] Therefore, it has great significance to isolate novel full-length cDNA clones of human, only few of which has been isolated. Especially, isolation of a novel cDNA clone encoding a secretory protein or membrane protein is desired since the protein itself would be useful as a medicine, and also the clones potentially include a gene associated with diseases. In addition, genes encoding proteins that are associated with signal transduction, glycoprotein, transcription, or diseases are expected to be useful as target molecules for therapy, or as medicines themselves. These genes form a gene group predicted to be strongly associated with diseases. Thus, identification of the full-length cDNA clones

encoding those proteins has great significance.

SUMMARY OF THE INVENTION

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[0010] An objective of the present invention is to provide a polynucleotide encoding a novel protein, a protein encoded by said polynucleotide, and novel usages of these.

[0011] The inventors have developed a method for efficiently cloning a human full-length cDNA that is predicted by the ATGpr etc. to be a full-length cDNA clone, from a full-length-enriched cDNA library that is synthesized by the oligocapping method. Then, the inventors determined the nucleotide sequence of the obtained cDNA clones from both 5'- and 3'- ends.

[0012] Furthermore, the inventors analyzed the obtained clones by the BLAST search of the databases, SwissProt (http://www.ebi.ac.uk/ebi_docsSwissProt_db/swisshome.html), GenBank (http://www.ncbi.nlm.nih.gov/web/GenBank), and UniGene (Human) (http://www.ncbi.nlm.nih.gov/UniGene).

[0013] The full-length cDNA clones of the present invention have high fullness ratio since these were obtained by the combination of (1) construction of a full-length-enriched cDNA library that is synthesized by the oligo-capping method, and (2) a system in which the full-length ratio is evaluated from the nucleotide sequence of the 5'-end (selection based on the ATGpr, previously removed complete sequences to ESTs). However, the primer of the present invention enables to obtain full-length cDNA easily without any specialized methods as in the described method.

Homology analysis in which the analysis is carried out against a not-full-length cDNA fragment to postulate the function of a protein encoded by said fragment, is being commonly performed.

However, since such analysis is based on the information of the fragment, it is not clear as to whether this fragment corresponds to a part that is functionally important in the protein. In other words, the reliability of the homology analysis based on the information of a fragment is doubtful, as information related to the structure of the whole protein is not available. However, the homology analysis of the present invention is conducted based on the information of a full-length cDNA comprising the whole coding region of the cDNA, and therefore, the homology of various portions of the protein can be analyzed. Hence, the reliability of the homology analysis has been dramatically improved in the present invention.

[0014] The inventors completed the invention by finding that it is possible to synthesize a novel full-length cDNA by using the combination of a primer that is designed based on the nucleotide sequence of the 5'-ends of the selected full-length cDNA clones and any of an oligo-dT primer or a 3'-primer that is designed based on the nucleotide sequence of the 3'-ends of the selected clones.

[0015] Thus, the present invention relates to primers described below, a method for synthesizing a polynucleotide using the primers, and polynucleotides obtained by the method.

[0016] First, the present invention relates to

(1) use of an oligonucleotide as a primer for synthesizing the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, or the complementary strand thereof, wherein said oligonucleotide is complementary to said polynucleotide or the complementary strand thereof and comprises at least 15 nucleotides;

(2) a primer set for synthesizing polynucleotides, the primer set comprising an oligo-dT primer and an oligonucleotide complementary to the complementary strand of the polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs: 1-5547 and SEQ ID NOs: 16111-16164, wherein said oligonucleotide comprises at least 15 nucleotides; and

(3) a primer set for synthesizing polynucleotides, the primer set comprising a combination of an oligonucleotide comprising a nucleotide sequence complementary to the complementary strand of the polynucleotide comprising a 5'-end nucleotide sequence and an oligonucleotide comprising a nucleotide sequence complementary to the polynucleotide comprising a 3'-end nucleotide sequence, wherein said oligonucleotides comprise at least 15 nucleotides and wherein said combination of 5'-end nucleotide sequence / 3'-end nucleotide sequence is selected from the combinations of 5'-end nucleotide sequence / 3'-end nucleotide sequence set forth in the SEQ ID NOs in Tables 1 and 2.

[0017] Tables 1 and 2 shows names of clones obtained in the examples described later, comprising the polynucleotide of the present invention (Table 1; 5547 clones, Table 2; 54 clones), names of nucleotide sequences at the 5'-end and 3'-end of the full-length cDNA, and their corresponding SEQ ID NOs. A blank indicates that the 3'-end sequence corresponding to the 5'-end sequence has not been determined for the same clone.

[0018] The SEQ ID NO of a 5'-end sequence is shown on the right side of the name of the 5'-end sequence, and the SEQ ID NO of a 3'-end sequence is shown on the right side of the name of the 3'-end sequence.

Table 1

				30 TD		922	270
name of	•	name of		EQ ID	name of		2 ID
clone		5'-end	of	5'-end	3'-end	of	3'-end
		sequence	se	equence	sequence	sec	quence
HEMBA10	00005	F-HEMBA10	00005	1	R-HEMBA100	0005	5548
HEMBA10	00012	F-HEMBA10	00012	2			
HEMBA10	00020	F-HEMBA10	00020	3			
HEMBA10	00030	F-HEMBA10	00030	4	R-HEMBA100	0030	5549
HEMBA10	00042	F-HEMBA10	00042	5	R-HEMBA100	0042	5550
HEMBA10	00046	F-HEMBA10	00046	6	R-HEMBA100	0046	5551
HEMBA10	00050	F-HEMBA10	00050	7	R-HEMBA100	0050	5552
HEMBA10	00076	F-HEMBA10	00076	8	R-HEMBA100	0076	5553
HEMBA10	000111	F-HEMBA10	00111	9	R-HEMBA100	0111	5554
HEMBA10	00129	F-HEMBA10	00129	10	R-HEMBA100	0129	5555
HEMBA10	000141	F-HEMBA10	00141	11	R-HEMBA100	0141	5556
HEMBA10	00150	F-HEMBA10	00150	12	R-HEMBA100	0150	5557
HEMBA10	000156	F-HEMBA10	00156	13	R-nnnnnnn	מממת	5558
HEMBA10	00158	F-HEMBA10	00158	14	R-HEMBA100	0158	5559
HEMBA1	000168	F-HEMBA10	00168	15	R-nnnnnnn	nnnn	5560

	HEMBA1000180	F-HEMBA1000180	16	R-HEMBA1000180	5561
	HEMBA1000185	F-HEMBA1000185	17	R-HEMBA1000185	5562
	HEMBA1000193	F-HEMBA1000193	18	R-HEMBA1000193	5563
5	HEMBA1000201	F-HEMBA1000201	19	R-HEMBA1000201	5564
	HEMBA1000213	F-HEMBA1000213	20	R-HEMBA1000213	5565
	HEMBA1000216	F-HEMBA1000216	21	R-HEMBA1000216	5566
	HEMBA1000227	F-HEMBA1000227	22	R-nnnnnnnnnnn	5567
10	HEMBA1000231	F-HEMBA1000231	23	R-HEMBA1000231	5568
	HEMBA1000243	F-HEMBA1000243	24	R-HEMBA1000243	5569
	HEMBA1000244	F-HEMBA1000244	25	R-HEMBA1000244	5570
	HEMBA1000251	F-HEMBA1000251	26	R-HEMBA1000251	5571
15	HEMBA1000264	F-HEMBA1000264	27	R-HEMBA1000264	5572
	HEMBA1000280	F-HEMBA1000280	28	R-nnnnnnnnnnn	5573
	HEMBA1000282	F-HEMBA1000282,	29	R-HEMBA1000282	5574
	HEMBA1000288	F-HEMBA1000288	30	R-HEMBA1000288	5575
20	HEMBA1000290	F-HEMBA1000290	31	R-HEMBA1000290	5576
	HEMBA1000302	F-HEMBA1000302	32	R-HEMBA1000302	5577
	HEMBA1000303	F-HEMBA1000303	33	R-nnnnnnnnnnn	5578
	HEMBA1000304	F-HEMBA1000304	34	R-nnnnnnnnnnn	5579
25	HEMBA1000307	F-HEMBA1000307	35	R-HEMBA1000307	5580
20	HEMBA1000327	F-HEMBA1000327	36		
	HEMBA1000333	F-HEMBA1000333	37	R-nnnnnnnnnn	5581
	HEMBA1000338	F-HEMBA1000338	38	R-HEMBA1000338	5582
30	HEMBA1000351	F-HEMBA1000351	39	R-HEMBA1000351	5583
30	HEMBA1000355	F-HEMBA1000355	40	R-HEMBA1000355	5584
	HEMBA1000356	F-HEMBA1000356	41		
	HEMBA1000357	F-HEMBA1000357	42	R-HEMBA1000357	5585
	HEMBA1000366	F-HEMBA1000366	43	R-HEMBA1000366	5586
35	HEMBA1000369	F-HEMBA1000369	44	R-HEMBA1000369	5587
	HEMBA1000376	F-HEMBA1000376	45	R-HEMBA1000376	5588
	HEMBA1000387	F-HEMBA1000387	46	R-HEMBA1000387	5589
	HEMBA1000390	F-HEMBA1000390	47	R-HEMBA1000390	5590
40	HEMBA1000392	F-HEMBA1000392	48	R-HEMBA1000392	5591
	HEMBA1000396	F-HEMBA1000396	49	R-HEMBA1000396	5592
	HEMBA1000411	F-HEMBA1000411	50	R-HEMBA1000411	5593
	HEMBA1000418	F-HEMBA1000418	51	R-HEMBA1000418	5594
45	HEMBA1000422	F-HEMBA1000422	52	R-HEMBA1000422	5595
	HEMBA1000428	F-HEMBA1000428	53	R-HEMBA1000428	5596
	HEMBA1000434	F-HEMBA1000434	5 4	R-HEMBA1000434	5597
	HEMBA1000442	F-HEMBA1000442	55	R-HEMBA1000442	5598
50	HEMBA1000456	F-HEMBA1000456	56	R-HEMBA1000456	5599
	HEMBA1000459	F-HEMBA1000459	57 50	R-HEMBA1000459	5600 5601
	HEMBA1000460	F-HEMBA1000460	58	R-HEMBA1000460	5601
	HEMBA1000464	F-HEMBA1000464	59	R-HEMBA1000464	5602
55	HEMBA1000469	F-HEMBA1000469	60	R-HEMBA1000469	5603
	HEMBA1000488	F-HEMBA1000488	61	R-HEMBA1000488	5604

	HEMBA1000490	F-HEMBA1000490	62	R-HEMBA1000490	5605
	HEMBA1000491	F-HEMBA1000491	63	R-HEMBA1000491	5606
	HEMBA1000501	F-HEMBA1000501	64		
5	HEMBA1000504	F-HEMBA1000504	65	R-HEMBA1000504	5607
	HEMBA1000505	F-HEMBA1000505	66	R-HEMBA1000505	5608
	HEMBA1000508	F-HEMBA1000508	67	R-HEMBA1000508	5609
	HEMBA1000518	F-HEMBA1000518	68	R-HEMBA1000518	5610
10	HEMBA1000519	F-HEMBA1000519	69	R-HEMBA1000519	5611
	HEMBA1000520	F-HEMBA1000520	70	R-HEMBA1000520	5612
	HEMBA1000523	F-HEMBA1000523	71	R-HEMBA1000523	5613
	HEMBA1000531	F-HEMBA1000531	72	R-HEMBA1000531	5614
15	HEMBA1000534	F-HEMBA1000534	73		
	HEMBA1000540	F-HEMBA1000540	74	R-HEMBA1000540	5615
	HEMBA1000542	F-HEMBA1000542,	75		
	HEMBA1000545	F-HEMBA1000545	76	R-HEMBA1000545	5616
20	HEMBA1000555	F-HEMBA1000555	77	R-nnnnnnnnnnn	5617
	HEMBA1000557	F-HEMBA1000557	78	R-HEMBA1000557	5618
	HEMBA1000561	F-HEMBA1000561	79	R-HEMBA1000561	5619
	HEMBA1000563	F-HEMBA1000563	80	R-HEMBA1000563	5620
25	HEMBA1000568	F-HEMBA1000568	81	R-HEMBA1000568	5621
25	HEMBA1000569	F-HEMBA1000569	82	R-nnnnnnnnnnn	5622
	HEMBA1000575	F-HEMBA1000575	83	R-HEMBA1000575	5623
	HEMBA1000588	F-HEMBA1000588	84	R-HEMBA1000588	5624
	HEMBA1000591	F-HEMBA1000591	85	R-HEMBA1000591	5625
30	HEMBA1000592	F-HEMBA1000592	86	R-HEMBA1000592	5626
	HEMBA1000594	F-HEMBA1000594	87	R-HEMBA1000594	5627
	HEMBA1000604	F-HEMBA1000604	88	R-HEMBA1000604	5628
	HEMBA1000608	F-HEMBA1000608	89	R-HEMBA1000608	5629
35	HEMBA1000622	F-HEMBA1000622	90	R-HEMBA1000622	5630
	HEMBA1000636	F-HEMBA1000636	91	R-HEMBA1000636	5631
	HEMBA1000637	F-HEMBA1000637	92	R-HEMBA1000637	5632
	HEMBA1000655	F-HEMBA1000655	93	R-HEMBA1000655	5633
40	HEMBA1000657	F-HEMBA1000657	94	R-HEMBA1000657	5634
	HEMBA1000662	F-HEMBA1000662	95	R-HEMBA1000662	5635
	HEMBA1000673	F-HEMBA1000673	96	R-HEMBA1000673	5636
	HEMBA1000682	F-HEMBA1000682	97	R-HEMBA1000682	5637
45	HEMBA1000686	F-HEMBA1000686	98	R-HEMBA1000686	5638
	HEMBA1000702	F-HEMBA1000702	99	R-HEMBA1000702	5639
	HEMBA1000705	F-HEMBA1000705	100	R-HEMBA1000705	5640
	HEMBA1000719	F-HEMBA1000719	101	R-HEMBA1000719	5641
50	HEMBA1000722	F-HEMBA1000722	102	R-HEMBA1000722	5642
	HEMBA1000726	F-HEMBA1000726	103	R-HEMBA1000726	5643
	HEMBA1000727	F-HEMBA1000727	104	R-HEMBA1000727	5644
	HEMBA1000747	F-HEMBA1000747	105	R-HEMBA1000747	5645
E E	HEMBA1000749	F-HEMBA1000749	106	R-HEMBA1000749	5646
55	HEMBA1000752	F-HEMBA1000752	107	R-HEMBA1000752	5647

	HEMBA1000769	F-HEMBA1000769	108	R-HEMBA1000769	5648
	HEMBA1000773	F-HEMBA1000773	109	R-HEMBA1000773	5649
	HEMBA1000774	F-HEMBA1000774	110	R-HEMBA1000774	5650
5	HEMBA1000791	F-HEMBA1000791	111	R-HEMBA1000791	5651
	HEMBA1000817	F-HEMBA1000817	112	R-HEMBA1000817	5652
	HEMBA1000822	F-HEMBA1000822	113	R-HEMBA1000822	5653
	HEMBA1000827	F-HEMBA1000827	114	R-HEMBA1000827	5654
10	HEMBA1000843	F-HEMBA1000843	115	R-HEMBA1000843	5655
	HEMBA1000851	F-HEMBA1000851	116	R-HEMBA1000851	5656
	HEMBA1000852	F-HEMBA1000852	117	R-HEMBA1000852	5657
	HEMBA1000867	F-HEMBA1000867	118	R-HEMBA1000867	5658
45	HEMBA1000869	F-HEMBA1000869	119	R-HEMBA1000869	5659
15	HEMBA1000870	F-HEMBA1000870	120	R-HEMBA1000870	5660
	HEMBA1000872	F-HEMBA1000872,	121	R-HEMBA1000872	5661
	HEMBA1000876	F-HEMBA1000876	122	R-HEMBA1000876	5662
	HEMBA1000908	F-HEMBA1000908	123	R-HEMBA1000908	5663
20	HEMBA1000910	F-HEMBA1000910	124	R-HEMBA1000910	5664
	HEMBA1000918	F-HEMBA1000918	125	R-HEMBA1000918	5665
	HEMBA1000919	F-HEMBA1000919	126	R-HEMBA1000919	5666
	HEMBA1000934	F-HEMBA1000934	127	R-HEMBA1000934	5667
25	HEMBA1000942	F-HEMBA1000942	128	R-HEMBA1000942	5668
	HEMBA1000943	F-HEMBA1000943	129	R-HEMBA1000943	5669
	HEMBA1000946	F-HEMBA1000946	130	R-HEMBA1000946	5670
	HEMBA1000960	F-HEMBA1000960	131	R-HEMBA1000960	5671
30	HEMBA1000968	F-HEMBA1000968	132	R-HEMBA1000968	5672
	HEMBA1000971	F-HEMBA1000971	133	R-HEMBA1000971	5673
	HEMBA1000972	F-HEMBA1000972	134	R-HEMBA1000972	5674
	HEMBA1000974	F-HEMBA1000974	135	R-HEMBA1000974	5675
35	HEMBA1000975	F-HEMBA1000975	136	R-HEMBA1000975	5676
	HEMBA1000985	F-HEMBA1000985	137	R-HEMBA1000985	5677
	HEMBA1000986	F-HEMBA1000986	138	R-HEMBA1000986	5678
	HEMBA1000991	F-HEMBA1000991	139	R-HEMBA1000991	5679
40	HEMBA1001007	F-HEMBA1001007	140	R-HEMBA1001007	5680
	HEMBA1001008	F-HEMBA1001008	141	R-HEMBA1001008	5681
	HEMBA1001009	F-HEMBA1001009	142	R-HEMBA1001009	5682
	HEMBA1001017	F-HEMBA1001017	143	R-HEMBA1001017	5683
4E	HEMBA1001019	F-HEMBA1001019	144	R-HEMBA1001019	5684
4 5	HEMBA1001020	F-HEMBA1001020	145	R-HEMBA1001020	5685
	HEMBA1001022	F-HEMBA1001022	146	R-HEMBA1001022	5686
	HEMBA1001024	F-HEMBA1001024	147	R-HEMBA1001024	5687
	HEMBA1001026	F-HEMBA1001026	148	R-HEMBA1001026	5688
50	HEMBA1001043	F-HEMBA1001043	149	R-nnnnnnnnnnn	5689
	HEMBA1001051	F-HEMBA1001051	150	R-HEMBA1001051	5690
	HEMBA1001052	F-HEMBA1001052	151	R-HEMBA1001052	5691
	HEMBA1001059	F-HEMBA1001059	152	•	
55	HEMBA1001060	F-HEMBA1001060	153	R-HEMBA1001060	5692

	·HEMBA1001071	F-HEMBA1001071	154	R-HEMBA1001071	5693
	HEMBA1001077	F-HEMBA1001077	155	R-HEMBA1001077	5694
	HEMBA1001080	F-HEMBA1001080	156	R-HEMBA1001080	5695
5	HEMBA1001085	F-HEMBA1001085	157	R-HEMBA1001085	5696
•	HEMBA1001088	F-HEMBA1001088	158	R-HEMBA1001088	5697
	HEMBA1001094	F-HEMBA1001094	159	R-HEMBA1001094	5698
	HEMBA1001099	F-HEMBA1001099	160	R-HEMBA1001099	5699
10	HEMBA1001109	F-HEMBA1001109	161	R-HEMBA1001109	5700
	HEMBA1001121	F-HEMBA1001121	162	R-HEMBA1001121	5701
	HEMBA1001122	F-HEMBA1001122	163	R-HEMBA1001122	5702
	HEMBA1001123	F-HEMBA1001123	164	R-HEMBA1001123	5703
15	HEMBA1001133	F-HEMBA1001133	165	R-HEMBA1001133	5704
.0	HEMBA1001137	F-HEMBA1001137	166	R-HEMBA1001137	5705
	HEMBA1001140	F-HEMBA1001140,	167	R-HEMBA1001140	5706
	HEMBA1001172	F-HEMBA1001172	168	R-HEMBA1001172	5707
	HEMBA1001174	F-HEMBA1001174	169	R-HEMBA1001174	5708
20	HEMBA1001197	F-HEMBA1001197	170	R-HEMBA1001197	5709
	HEMBA1001208	F-HEMBA1001208	171	R-HEMBA1001208	5710
	HEMBA1001213	F-HEMBA1001213	172		
	HEMBA1001226	F-HEMBA1001226	173	R-HEMBA1001226	5711
25	HEMBA1001235	F-HEMBA1001235	174	R-HEMBA1001235	5712
	HEMBA1001247	F-HEMBA1001247	175	R-HEMBA1001247	5713
	HEMBA1001257	F-HEMBA1001257	176	R-HEMBA1001257	5714
	HEMBA1001265	F-HEMBA1001265	177	R-HEMBA1001265	5715
30	HEMBA1001281	F-HEMBA1001281	178	R-nnnnnnnnnnn	5716
	HEMBA1001286	F-HEMBA1001286	179	R-HEMBA1001286	5717
	HEMBA1001289	F-HEMBA1001289	180	R-HEMBA1001289	5718
	HEMBA1001294	F-HEMBA1001294	181	R-HEMBA1001294	5719
35	HEMBA1001299	F-HEMBA1001299	182	R-HEMBA1001299	5720
	HEMBA1001302	F-HEMBA1001302	183	R-HEMBA1001302	5721
	HEMBA1001303	F-HEMBA1001303	184	R-HEMBA1001303	5722
	HEMBA1001310	F-HEMBA1001310	185	R-HEMBA1001310	5723
40	HEMBA1001319	F-HEMBA1001319	186	R-HEMBA1001319	5724
	HEMBA1001323	F-HEMBA1001323	187	R-HEMBA1001323	5725
	HEMBA1001326	F-HEMBA1001326	188	R-HEMBA1001326	5726
	HEMBA1001327	F-HEMBA1001327	189	R-HEMBA1001327	5727
45	HEMBA1001330	F-HEMBA1001330	190	R-HEMBA1001330	5728
45	HEMBA1001351	F-HEMBA1001351	191	R-HEMBA1001351	5729
	HEMBA1001361	F-HEMBA1001361	192	R-HEMBA1001361	5730
	HEMBA1001375	F-HEMBA1001375	193	R-HEMBA1001375	5731
	HEMBA1001377	F-HEMBA1001377	194	R-HEMBA1001377	5732
50	HEMBA1001383	F-HEMBA1001383	195	R-HEMBA1001383	5733
	HEMBA1001387	F-HEMBA1001387	196	R-HEMBA1001387	5734
	HEMBA1001388	F-HEMBA1001388	197	R-HEMBA1001388	5735
	HEMBA1001391	F-HEMBA1001391	198	R-HEMBA1001391	5736
55	HEMBA1001398	F-HEMBA1001398	199	R-HEMBA1001398	5737

	HEMBA1001405	F-HEMBA1001405	200	R-HEMBA1001405	5738
	HEMBA1001407	F-HEMBA1001407	201	R-HEMBA1001407	5739
	HEMBA1001411	F-HEMBA1001411	202	R-HEMBA1001411	5740
5	HEMBA1001413	F-HEMBA1001413	203	R-HEMBA1001413	5741
	HEMBA1001415	F-HEMBA1001415	204	R-HEMBA1001415	5742
	HEMBA1001432	F-HEMBA1001432	205	R-HEMBA1001432	5743
	HEMBA1001433	F-HEMBA1001433	206	R-HEMBA1001433	5744
10	HEMBA1001435	F-HEMBA1001435	207	R-HEMBA1001435	5745
	HEMBA1001442	F-HEMBA1001442	208	R-HEMBA1001442	5746
	HEMBA1001446	F-HEMBA1001446	209	R-HEMBA1001446	5747
	HEMBA1001450	F-HEMBA1001450	210	R-HEMBA1001450	5748
15	HEMBA1001454	F-HEMBA1001454	211	R-HEMBA1001454	5749
	HEMBA1001455	F-HEMBA1001455	212	R-HEMBA1001455	5750
	HEMBA1001463	F-HEMBA1001463,	213	R-HEMBA1001463	5751
	HEMBA1001476	F-HEMBA1001476	214	R-HEMBA1001476	5752
20	HEMBA1001478	F-HEMBA1001478	215	R-HEMBA1001478	5753
20	HEMBA1001497	F-HEMBA1001497	216	R-HEMBA1001497	5754
	HEMBA1001510	F-HEMBA1001510	217	R-HEMBA1001510	5755
	HEMBA1001515	F-HEMBA1001515	218	R-HEMBA1001515	5756
	HEMBA1001517	F-HEMBA1001517	219	R-HEMBA1001517	5757
25	HEMBA1001522	F-HEMBA1001522	220	R-HEMBA1001522	5758
	HEMBA1001526	F-HEMBA1001526	221	R-HEMBA1001526	5759
	HEMBA1001533	F-HEMBA1001533	222	R-HEMBA1001533	5760
	HEMBA1001557	F-HEMBA1001557	223	R-HEMBA1001557	5761
30	HEMBA1001566	F-HEMBA1001566	224	R-HEMBA1001566	5762
	HEMBA1001569	F-HEMBA1001569	225	R-HEMBA1001569	5763
	HEMBA1001570	F-HEMBA1001570	226	R-HEMBA1001570	5764
	HEMBA1001579	F-HEMBA1001579	227	R-HEMBA1001579	5765
35	HEMBA1001581	F-HEMBA1001581	228	R-HEMBA1001581	5766
	HEMBA1001585	F-HEMBA1001585	229	R-HEMBA1001585	5767
	HEMBA1001589	F-HEMBA1001589	230	R-HEMBA1001589	5768
	HEMBA1001595	F-HEMBA1001595	231	R-HEMBA1001595	5769
40	HEMBA1001608	F-HEMBA1001608	232	R-HEMBA1001608	5770
	HEMBA1001620	F-HEMBA1001620	233	R-HEMBA1001620	5771
	HEMBA1001635	F-HEMBA1001635	234	R-nnnnnnnnnnnn	5772
	HEMBA1001636	F-HEMBA1001636	235	R-HEMBA1001636	5773
45	HEMBA1001640	F-HEMBA1001640	236	R-HEMBA1001640	5774
	HEMBA1001647	F-HEMBA1001647	237		
	HEMBA1001651	F-HEMBA1001651	238	R-nnnnnnnnnnn	5775
	HEMBA1001655	F-HEMBA1001655	239	R-HEMBA1001655	5776
50	HEMBA1001658	F-HEMBA1001658	240	R-HEMBA1001658	5777
50	HEMBA1001661	F-HEMBA1001661	241	R-HEMBA1001661	5778
	HEMBA1001672	F-HEMBA1001672	242	R-HEMBA1001672	5779
	HEMBA1001675	F-HEMBA1001675	243	R-HEMBA1001675	5780
	HEMBA1001678	F-HEMBA1001678	244	R-HEMBA1001678	5781
55	HEMBA1001681	F-HEMBA1001681	245	R-HEMBA1001681	5782

	HEMBA1001702	F-HEMBA1001702	246	R-HEMBA1001702	5783
	HEMBA1001709	F-HEMBA1001709	247	R-HEMBA1001709	5784
	HEMBA1001711	F-HEMBA1001711	248	R-HEMBA1001711	5785
5	HEMBA1001712	F-HEMBA1001712	249	R-HEMBA1001712	5786
	HEMBA1001714	F-HEMBA1001714	250	R-HEMBA1001714	5787
	HEMBA1001718	F-HEMBA1001718	251	R-HEMBA1001718	5788
	HEMBA1001723	F-HEMBA1001723	252	R-HEMBA1001723	5789
10	HEMBA1001731	F-HEMBA1001731	253	R-HEMBA1001731	5790
	HEMBA1001734	F-HEMBA1001734	254	R-HEMBA1001734	5791
	HEMBA1001744	F-HEMBA1001744	255	R-HEMBA1001744	5792
	HEMBA1001745	F-HEMBA1001745	256	R-HEMBA1001745	5793
15	HEMBA1001746	F-HEMBA1001746	257	R-HEMBA1001746	5794
	HEMBA1001761	F-HEMBA1001761	258	R-HEMBA1001761	5795
	HEMBA1001781	F-HEMBA1001781,	259	R-HEMBA1001781	5796
	HEMBA1001784	F-HEMBA1001784	260	R-HEMBA1001784	5797
20	HEMBA1001791	F-HEMBA1001791	261	R-HEMBA1001791	5798
	HEMBA1001800	F-HEMBA1001800	262	R-HEMBA1001800	5799
	HEMBA1001803	F-HEMBA1001803	263	R-HEMBA1001803	5800
	HEMBA1001804	F-HEMBA1001804	264	R-nnnnnnnnnnn	5801
25	HEMBA1001808	F-HEMBA1001808	265	R-HEMBA1001808	5802
25	HEMBA1001809	F-HEMBA1001809	266	R-HEMBA1001809	5803
	HEMBA1001815	F-HEMBA1001815	267	R-HEMBA1001815	5804
	HEMBA1001819	F-HEMBA1001819	268	R-HEMBA1001819	5805
	HEMBA1001820	F-HEMBA1001820	269	R-HEMBA1001820	5806
30	HEMBA1001822	F-HEMBA1001822	270	R-nnnnnnnnnnn	5807
	HEMBA1001824	F-HEMBA1001824	271	R-HEMBA1001824	5808
	HEMBA1001835	F-HEMBA1001835	272	R-HEMBA1001835	5809
	HEMBA1001844	F-HEMBA1001844	273	R-HEMBA1001844	5810
35	HEMBA1001847 HEMBA1001861	F-HEMBA1001847	274	R-HEMBA1001847	5811 5812
	HEMBA1001864	F-HEMBA1001861 F-HEMBA1001864	275 276	R-HEMBA1001861 R-HEMBA1001864	5812
	HEMBA1001866	F-HEMBA1001866	277	R-HEMBA1001866	5814
	HEMBA1001869	F-HEMBA1001869	278	R-nnnnnnnnnn	5815
40	HEMBA1001888	F-HEMBA1001888	279	R-HEMBA1001888	5816
	HEMBA1001896	F-HEMBA1001896	280	R-HEMBA1001886	5817
	HEMBA1001910	F-HEMBA1001910	281	R-HEMBA1001910	5818
	HEMBA1001912	F-HEMBA1001912	282	R-HEMBA1001912	5819
45	HEMBA1001913	F-HEMBA1001913	283	R-HEMBA1001913	5820
	HEMBA1001915	F-HEMBA1001915	284	R-HEMBA1001915	5821
	HEMBA1001918	F-HEMBA1001918	285	R-HEMBA1001918	5822
	HEMBA1001921	F-HEMBA1001921	286	R-HEMBA1001921	5823
50	HEMBA1001939	F-HEMBA1001939	287	R-HEMBA1001939	5824
	HEMBA1001940	F-HEMBA1001940	288	R-HEMBA1001940	5825
	HEMBA1001942	F-HEMBA1001942	289	R-HEMBA1001942	5826
	HEMBA1001945	F-HEMBA1001945	290	R-HEMBA1001945	5827
55	HEMBA1001950	F-HEMBA1001950	291	R-HEMBA1001950	5828

	HEMBA1001960	F-HEMBA1001960	292	R-HEMBA1001960	5829
	HEMBA1001962	F-HEMBA1001962	293	R-HEMBA1001962	5830
	HEMBA1001964	F-HEMBA1001964	294	R-HEMBA1001964	5831
5	HEMBA1001967	F-HEMBA1001967	295	R-HEMBA1001967	5832
	HEMBA1001979	F-HEMBA1001979	296	R-HEMBA1001979	5833
	HEMBA1001987	F-HEMBA1001987	297	R-HEMBA1001987	5834
	HEMBA1001991	F-HEMBA1001991	298	R-HEMBA1001991	5835
10	HEMBA1002003	F-HEMBA1002003	299	R-HEMBA1002003	5836
	HEMBA1002008	F-HEMBA1002008	300	R-HEMBA1002008	5837
	HEMBA1002018	F-HEMBA1002018	301	R-HEMBA1002018	5838
	HEMBA1002022	F-HEMBA1002022	302	R-HEMBA1002022	5839
45	HEMBA1002035	F-HEMBA1002035	303	R-HEMBA1002035	5840
15	HEMBA1002039	F-HEMBA1002039	304	R-HEMBA1002039	5841
	HEMBA1002049	F-HEMBA1002049	305	R-HEMBA1002049	5842
	HEMBA1002084	F-HEMBA1002084	306	R-HEMBA1002084	5843
	HEMBA1002092	F-HEMBA1002092	307	R-HEMBA1002092	5844
20	HEMBA1002100	F-HEMBA1002100	308	R-HEMBA1002100	5845
	HEMBA1002102	F-HEMBA1002102	309	R-HEMBA1002102	5846
	HEMBA1002113	F-HEMBA1002113	310	R-HEMBA1002113	5847
	HEMBA1002119	F-HEMBA1002119	311	R-HEMBA1002119	5848
25	HEMBA1002125	F-HEMBA1002125	312	R-HEMBA1002125	5849
	HEMBA1002139	F-HEMBA1002139	313	R-HEMBA1002139	5850
	HEMBA1002144	F-HEMBA1002144	314	R-HEMBA1002144	5851
	HEMBA1002150	F-HEMBA1002150	315	R-HEMBA1002150	5852
30	HEMBA1002151	F-HEMBA1002151	316	R-HEMBA1002151	5853
	HEMBA1002153	F-HEMBA1002153	317	R-HEMBA1002153	5854
	HEMBA1002160	F-HEMBA1002160	318	R-HEMBA1002160	5855
	HEMBA1002161	F-HEMBA1002161	319	R-HEMBA1002161	5856
35	HEMBA1002162	F-HEMBA1002162	320	R-HEMBA1002162	5857
	HEMBA1002166	F-HEMBA1002166	321	R-HEMBA1002166	5858
	HEMBA1002177	F-HEMBA1002177	322	R-HEMBA1002177	5859
	HEMBA1002185	F-HEMBA1002185	323	R-HEMBA1002185	5860
40	HEMBA1002189	F-HEMBA1002189	324	R-HEMBA1002189	5861
••	HEMBA1002191	F-HEMBA1002191	325	R-HEMBA1002191	5862
	HEMBA1002199	F-HEMBA1002199	326	R-HEMBA1002199	5863
	HEMBA1002204	F-HEMBA1002204	327	R-HEMBA1002204	5864
	HEMBA1002212	F-HEMBA1002212	328	R-HEMBA1002212	5865
45	HEMBA1002215	F-HEMBA1002215	329	R-HEMBA1002215	5866
	HEMBA1002226	F-HEMBA1002226	330	R-HEMBA1002226	5867
	HEMBA1002229	F-HEMBA1002229	331	R-HEMBA1002229	5868
	HEMBA1002237	F-HEMBA1002237	332	R-HEMBA1002237	5869
50	HEMBA1002241	F-HEMBA1002241	333	- 11mm	
	HEMBA1002253	F-HEMBA1002253	334	R-HEMBA1002253	5870
	HEMBA1002257	F-HEMBA1002257	335	R-HEMBA1002257	5871
	HEMBA1002265	F-HEMBA1002265	336	D 100001-00005	5050
55	HEMBA1002267	F-HEMBA1002267	337	R-HEMBA1002267	5872

	HEMBA1002270.	F-HEMBA1002270	338	R-HEMBA1002270	5873
	HEMBA1002321	F-HEMBA1002321	339	R-HEMBA1002321	5874
		F-HEMBA1002328	340	R-HEMBA1002328	5875
5	HEMBA1002337	F-HEMBA1002337	341	R-HEMBA1002337	5876
	HEMBA1002331	F-HEMBA1002341	342	R-HEMBA1002341	5877
	HEMBA1002348	F-HEMBA1002348	343	R-HEMBA1002348	5878
	HEMBA1002349	F-HEMBA1002349	344	R-HEMBA1002349	5879
40			345	R-nnnnnnnnnnn	5880
10	HEMBA1002363	F-HEMBA1002363 F-HEMBA1002381	345 346	R-HEMBA1002381	5881
	HEMBA1002381 HEMBA1002389	•		R-HEMBA1002389	5882
		F-HEMBA1002389	347		5883
	HEMBA1002417	F-HEMBA1002417	348	R-HEMBA1002417	
15	HEMBA1002419	F-HEMBA1002419	349	R-HEMBA1002419	5884
	HEMBA1002430	F-HEMBA1002430	350	R-HEMBA1002430	5885
	HEMBA1002439	F-HEMBA1002439,	351	R-HEMBA1002439	5886
	HEMBA1002458	F-HEMBA1002458	352	R-HEMBA1002458	5887
20	HEMBA1002460	F-HEMBA1002460	353	R-HEMBA1002460	5888
	HEMBA1002462	F-HEMBA1002462	354	R-HEMBA1002462	5889
	HEMBA1002469	F-HEMBA1002469	355	_	5000
	HEMBA1002475	F-HEMBA1002475	356	R-nnnnnnnnnnn	5890
25	HEMBA1002477	F-HEMBA1002477	357	R-HEMBA1002477	5891
25	HEMBA1002486	F-HEMBA1002486	358	R-HEMBA1002486	5892
	HEMBA1002495	F-HEMBA1002495	359	R-HEMBA1002495	5893
	HEMBA1002498	F-HEMBA1002498	360	R-HEMBA1002498	5894
	HEMBA1002503	F-HEMBA1002503	361	R-HEMBA1002503	5895
30	HEMBA1002508	F-HEMBA1002508	362	R-HEMBA1002508	5896
	HEMBA1002513	F-HEMBA1002513	363	R-nnnnnnnnnnn	5897
	HEMBA1002515	F-HEMBA1002515	364	R-HEMBA1002515	5898
	HEMBA1002538	F-HEMBA1002538	365	R-HEMBA1002538	5899
35	HEMBA1002542	F-HEMBA1002542	366	R-HEMBA1002542	5900
	HEMBA1002547	F-HEMBA1002547	367	R-HEMBA1002547	5901
	HEMBA1002552	F-HEMBA1002552	368	R-HEMBA1002552	5902
	HEMBA1002555	F-HEMBA1002555	369	R-HEMBA1002555	5903
40	HEMBA1002558	F-HEMBA1002558	370	R-HEMBA1002558	5904
	HEMBA1002561	F-HEMBA1002561	371	R-HEMBA1002561	5905
	HEMBA1002569	F-HEMBA1002569	372	R-nnnnnnnnnnn	5906
	HEMBA1002583	F-HEMBA1002583	373	R-HEMBA1002583	5907
45	HEMBA1002590	F-HEMBA1002590	374	R-HEMBA1002590	5908
40	HEMBA1002592	F-HEMBA1002592	375	R-HEMBA1002592	5909
	HEMBA1002609	F-HEMBA1002609	376	D IEMP41000001	C010
	HEMBA1002621	F-HEMBA1002621	377	R-HEMBA1002621	5910
	HEMBA1002624	F-HEMBA1002624	378	R-HEMBA1002624	5911
50	HEMBA1002628	F-HEMBA1002628	379	R-HEMBA1002628	5912
	HEMBA1002629	F-HEMBA1002629	380	R-HEMBA1002629	5913
	HEMBA1002645	F-HEMBA1002645	381	R-HEMBA1002645	5914 5915
	HEMBA1002651	F-HEMBA1002651	382	R-HEMBA1002651	
55	HEMBA1002659	F-HEMBA1002659	383	R-HEMBA1002659	5916

	HEMBA1002661	F-HEMBA1002661	384	R-HEMBA1002661	5917
	HEMBA1002666	F-HEMBA1002666	385	R-HEMBA1002666	5918
	HEMBA1002678	F-HEMBA1002678	386	R-HEMBA1002678	5919
5	HEMBA1002679	F-HEMBA1002679	387	R-nnnnnnnnnnn	5920
	HEMBA1002688	F-HEMBA1002688	388	R-HEMBA1002688	5921
	HEMBA1002696	F-HEMBA1002696	389	R-HEMBA1002696	5922
	HEMBA1002703	F-HEMBA1002703	390		
10	HEMBA1002712	F-HEMBA1002712	391	R-HEMBA1002712	5923
	HEMBA1002716	F-HEMBA1002716	392	R-HEMBA1002716	5924
	HEMBA1002728	F-HEMBA1002728	393	R-HEMBA1002728	5925
	HEMBA1002730	F-HEMBA1002730	394	R-HEMBA1002730	5926
15	HEMBA1002742	F-HEMBA1002742	395	R-HEMBA1002742	5927
70	HEMBA1002746	F-HEMBA1002746	396	R-HEMBA1002746	5928
	HEMBA1002748	F-HEMBA1002748,	397	R-HEMBA1002748	5929
	HEMBA1002750	F-HEMBA1002750	398	R-HEMBA1002750	5930
	HEMBA1002768	F-HEMBA1002768	399	R-HEMBA1002768	5931
20	HEMBA1002770	F-HEMBA1002770	400	R-HEMBA1002770	5932
	HEMBA1002777	F-HEMBA1002777	401	R-HEMBA1002777	5933
	HEMBA1002779	F-HEMBA1002779	402	R-HEMBA1002779	5934
	HEMBA1002780	F-HEMBA1002780	403	R-HEMBA1002780	5935
25	HEMBA1002794	F-HEMBA1002794	404	R-HEMBA1002794	5936
	HEMBA1002801	F-HEMBA1002801	405	R-HEMBA1002801	5937
	HEMBA1002810	F-HEMBA1002810	406	R-HEMBA1002810	5938
	HEMBA1002816	F-HEMBA1002816	407	R-HEMBA1002816	5939
30	HEMBA1002818	F-HEMBA1002818	408		
	HEMBA1002826	F-HEMBA1002826	409	R-HEMBA1002826	5940
	HEMBA1002833	F-HEMBA1002833	410	R-HEMBA1002833	5941
	HEMBA1002850	F-HEMBA1002850	411	R-HEMBA1002850	5942
35	HEMBA1002863	F-HEMBA1002863	412	R-HEMBA1002863	5943
	HEMBA1002876	F-HEMBA1002876	413	R-HEMBA1002876	5944
	HEMBA1002886	F-HEMBA1002886	414	R-HEMBA1002886	5945
	HEMBA1002896	F-HEMBA1002896	415	R-HEMBA1002896	5946
40	HEMBA1002921	F-HEMBA1002921	416	R-HEMBA1002921	5947
40	HEMBA1002924	F-HEMBA1002924	417	R-HEMBA1002924	5948
	HEMBA1002934	F-HEMBA1002934	418	R-HEMBA1002934	5949
	HEMBA1002935	F-HEMBA1002935	419	R-HEMBA1002935	5950
	HEMBA1002937	F-HEMBA1002937	420	R-HEMBA1002937	5951
45	HEMBA1002939	F-HEMBA1002939	421	R-HEMBA1002939	5952
	HEMBA1002944	F-HEMBA1002944	422	R-HEMBA1002944	5953
	HEMBA1002951	F-HEMBA1002951	423	R-HEMBA1002951	5954
	HEMBA1002954	F-HEMBA1002954	424	R-HEMBA1002954	5955
50	HEMBA1002968	F-HEMBA1002968	425	R-HEMBA1002968	5956
	HEMBA1002970	F-HEMBA1002970	426	R-HEMBA1002970	5957
	HEMBA1002971	F-HEMBA1002971	427	R-HEMBA1002971	5958
	HEMBA1002973	F-HEMBA1002973	428	R-HEMBA1002973	5959
55	HEMBA1002997	F-HEMBA1002997	429	R-nnnnnnnnnnn	5960

	HEMBA1002999	F-HEMBA1002999	430	R-HEMBA1002999	5961
	HEMBA1003021	F-HEMBA1003021	431	R-HEMBA1003021	5962
	HEMBA1003033	F-HEMBA1003033	432	R-HEMBA1003033	5963
5	HEMBA1003034	F-HEMBA1003034	433	R-HEMBA1003034	5964
	HEMBA1003035	F-HEMBA1003035	434	R-HEMBA1003035	5965
	HEMBA1003037	F-HEMBA1003037	435	R-HEMBA1003037	5966
	HEMBA1003041	F-HEMBA1003041	436	R-HEMBA1003041	5967
10	HEMBA1003041	F-HEMBA1003041	437	R-HEMBA1003046	5968
	HEMBA1003064	F-HEMBA1003064	438	R-HEMBA1003064	5969
	HEMBA1003067	F-HEMBA1003067	439	R-HEMBA1003067	5970
				R-HEMBA1003071	5971
	HEMBA1003071	F-HEMBA1003071	440	•	5971
15	HEMBA1003077	F-HEMBA1003077	441	R-HEMBA1003077	
	HEMBA1003078	F-HEMBA1003078	442	R-HEMBA1003078	5973
	HEMBA1003079	F-HEMBA1003079,	443	R-HEMBA1003079	5974
	HEMBA1003083	F-HEMBA1003083	444	R-HEMBA1003083	5975
20	HEMBA1003086	F-HEMBA1003086	445	R-HEMBA1003086	5976
	HEMBA1003096	F-HEMBA1003096	446	R-HEMBA1003096	5977
	HEMBA1003098	F-HEMBA1003098	447	R-HEMBA1003098	5978
	HEMBA1003117	F-HEMBA1003117	448	R-HEMBA1003117	5979
25	HEMBA1003129	F-HEMBA1003129	449	R-HEMBA1003129	5980
25	HEMBA1003133	F-HEMBA1003133	450	R-HEMBA1003133	5981
	HEMBA1003136	F-HEMBA1003136	451	R-HEMBA1003136	5982
	HEMBA1003142	F-HEMBA1003142	452	R-HEMBA1003142	5983
	HEMBA1003148	F-HEMBA1003148	453	R-HEMBA1003148	5984
30	HEMBA1003166	F-HEMBA1003166	454	R-HEMBA1003166	5985
	HEMBA1003175	F-HEMBA1003175	455	R-HEMBA1003175	5986
	HEMBA1003179	F-HEMBA1003179	456		
	HEMBA1003197	F-HEMBA1003197	457	R-HEMBA1003197	5987
35	HEMBA1003199	F-HEMBA1003199	458	R-HEMBA1003199	5988
	HEMBA1003202	F-HEMBA1003202	459	R-HEMBA1003202	5989
	HEMBA1003204	F-HEMBA1003204	460	R-HEMBA1003204	5990
	HEMBA1003212	F-HEMBA1003212	461	R-HEMBA1003212	5991
40	HEMBA1003220	F-HEMBA1003220	462	R-HEMBA1003220	5992
40	HEMBA1003222	F-HEMBA1003222	463	R-HEMBA1003222	5993
	HEMBA1003229	F-HEMBA1003229	464	R-HEMBA1003229	5994
	HEMBA1003235	F-HEMBA1003235	465	R-HEMBA1003235	5995
	HEMBA1003250	F-HEMBA1003250	466	R-HEMBA1003250	5996
45	HEMBA1003257	F-HEMBA1003257	467	R-HEMBA1003257	5997
	HEMBA1003273	F-HEMBA1003273	468	R-HEMBA1003273	5998
	HEMBA1003276	F-HEMBA1003276	469	R-HEMBA1003276	5999
	HEMBA1003278	F-HEMBA1003278	470	R-HEMBA1003278	6000
50	HEMBA1003281	F-HEMBA1003281	471	R-HEMBA1003281	6001
	HEMBA1003286	F-HEMBA1003286	472		
	HEMBA1003291	F-HEMBA1003291	473	R-HEMBA1003291	6002
	HEMBA1003296	F-HEMBA1003296	474	R-HEMBA1003296	6003
55	HEMBA1003304	F-HEMBA1003304	475	R-HEMBA1003304	6004
55					-

	HEMBA1003309	F-HEMBA1003309	476	R-HEMBA1003309	6005
	HEMBA1003314	F-HEMBA1003314	477	R-HEMBA1003314	6006
	HEMBA1003322	F-HEMBA1003322	478	R-HEMBA1003322	6007
5	HEMBA1003327	F-HEMBA1003327	479	R-HEMBA1003327	6008
	HEMBA1003328	F-HEMBA1003328	480	R-HEMBA1003328	6009
	HEMBA1003330	F-HEMBA1003330	481	R-HEMBA1003330	6010
	HEMBA1003348	F-HEMBA1003348	482	R-HEMBA1003348	6011
10	HEMBA1003369	F-HEMBA1003369	483	R-HEMBA1003369	6012
	HEMBA1003370	F-HEMBA1003370	484	R-HEMBA1003370	6013
	HEMBA1003373	F-HEMBA1003373	485	R-HEMBA1003373	6014
	HEMBA1003376	F-HEMBA1003376	486	R-HEMBA1003376	6015
15	HEMBA1003380	F-HEMBA1003380	487	R-HEMBA1003380	6016
10	HEMBA1003384	F-HEMBA1003384	488	R-HEMBA1003384	6017
	HEMBA1003395	F-HEMBA1003395,	489	R-HEMBA1003395	6018
	HEMBA1003402	F-HEMBA1003402	490	R-HEMBA1003402	6019
	HEMBA1003403	F-HEMBA1003403	491		0010
20	HEMBA1003408	F-HEMBA1003408	492	R-nnnnnnnnnnn	6020
	HEMBA1003417	F-HEMBA1003417	493	R-HEMBA1003417	6021
	HEMBA1003418	F-HEMBA1003418	494	R-HEMBA1003418	6022
	HEMBA1003433	F-HEMBA1003433	495	R-HEMBA1003433	6023
25	HEMBA1003447	F-HEMBA1003447	496		0020
	HEMBA1003461	F-HEMBA1003461	497	R-HEMBA1003461	6024
	HEMBA1003463	F-HEMBA1003463	498	R-HEMBA1003463	6025
	HEMBA1003480	F-HEMBA1003480	499	R-HEMBA1003480	6026
30	HEMBA1003528	F-HEMBA1003528	500	R-HEMBA1003528	6027
	HEMBA1003531	F-HEMBA1003531	501	R-HEMBA1003531	6028
	HEMBA1003538	F-HEMBA1003538	502	R-HEMBA1003538	6029
	HEMBA1003545	F-HEMBA1003545	503	R-HEMBA1003545	6030
35	HEMBA1003548	F-HEMBA1003548	504	R-HEMBA1003548	6031
	HEMBA1003555	F-HEMBA1003555	505	R-HEMBA1003555	6032
	HEMBA1003556	F-HEMBA1003556	506	R-HEMBA1003556	6033
	HEMBA1003560	F-HEMBA1003560	507	R-HEMBA1003560	6034
40	HEMBA1003568	F-HEMBA1003568	508	R-HEMBA1003568	6035
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	HEMBA1003571	F-HEMBA1003571	510	R-HEMBA1003571	6037
	HEMBA1003579	F-HEMBA1003579	511	R-HEMBA1003579	6038
	HEMBA1003581	F-HEMBA1003581	512	R-HEMBA1003581	6039
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	HEMBA1003595	F-HEMBA1003595	514	R-HEMBA1003595	6041
	HEMBA1003597	F-HEMBA1003597	515	R-HEMBA1003597	6042
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50	HEMBA1003615	F-HEMBA1003615	517	R-HEMBA1003615	6044
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	HEMBA1003621	F-HEMBA1003621	519	R-HEMBA1003621	6046
	HEMBA1003622	F-HEMBA1003622	520	R-HEMBA1003622	6047
55	HEMBA1003630	F-HEMBA1003630	521	R-HEMBA1003630	6048

	HEMBA1003637	F-HEMBA1003637	522	R-HEMBA1003637	6049
	HEMBA1003640	F-HEMBA1003640	523	R-HEMBA1003640	6050
	HEMBA1003645	F-HEMBA1003645	524	R-HEMBA1003645	6051
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	HEMBA1003656	F-HEMBA1003656	526	R-HEMBA1003656	6053
	HEMBA1003662	F-HEMBA1003662	527	R-HEMBA1003662	6054
	HEMBA1003667	F-HEMBA1003667	528	R-HEMBA1003667	6055
10	HEMBA1003679	F-HEMBA1003679	529	R-HEMBA1003679	6056
	HEMBA1003680	F-HEMBA1003680	530	R-HEMBA1003680	6057
	HEMBA1003684	F-HEMBA1003684	531	R-HEMBA1003684	6058
	HEMBA1003690	F-HEMBA1003690	532	R-HEMBA1003690	6059
	HEMBA1003692	F-HEMBA1003692	533	R-HEMBA1003692	6060
15	HEMBA1003711	F-HEMBA1003711	534	R-HEMBA1003711	6061
	HEMBA1003714	F-HEMBA1003714,	535	R-HEMBA1003714	6062
	HEMBA1003715	F-HEMBA1003715	536	R-HEMBA1003715	6063
	HEMBA1003720	F-HEMBA1003720	537	R-HEMBA1003720	6064
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	HEMBA1003729	F-HEMBA1003729	539	R-HEMBA1003729	6066
	HEMBA1003733	F-HEMBA1003733	540	R-HEMBA1003733	6067
	HEMBA1003742	F-HEMBA1003742	541	R-HEMBA1003742	6068
25	HEMBA1003758	F-HEMBA1003758	542	R-HEMBA1003758	6069
	HEMBA1003760	F-HEMBA1003760	543	R-HEMBA1003760	6070
	HEMBA1003773	F-HEMBA1003773	544	R-HEMBA1003773	6071
	HEMBA1003783	F-HEMBA1003783	545	R-HEMBA1003783	6072
30	HEMBA1003784	F-HEMBA1003784	546	R-HEMBA1003784	6073
30	HEMBA1003799	F-HEMBA1003799	547	R-HEMBA1003799	6074
	HEMBA1003803	F-HEMBA1003803	548	R-HEMBA1003803	6075
	HEMBA1003804	F-HEMBA1003804	549	R-HEMBA1003804	6076
	HEMBA1003805	F-HEMBA1003805	550	R-HEMBA1003805	6077
35	HEMBA1003807	F-HEMBA1003807	551	R-HEMBA1003807	6078
	HEMBA1003827	F-HEMBA1003827	552		
	HEMBA1003836	F-HEMBA1003836	553	R-HEMBA1003836	6079
	HEMBA1003838	F-HEMBA1003838	554	R-HEMBA1003838	6080
40	HEMBA1003856	F-HEMBA1003856	555	R-HEMBA1003856	6081
	HEMBA1003864	F-HEMBA1003864	556	R-HEMBA1003864	6082
	HEMBA1003866	F-HEMBA1003866	557	R-HEMBA1003866	6083
	HEMBA1003879	F-HEMBA1003879	558	R-HEMBA1003879	6084
45	HEMBA1003880	F-HEMBA1003880	559	R-HEMBA1003880	6085
	HEMBA1003885	F-HEMBA1003885	560	R-HEMBA1003885	6086
	HEMBA1003893	F-HEMBA1003893	561	R-HEMBA1003893	6087
	HEMBA1003902	F-HEMBA1003902	562	R-HEMBA1003902	6088
50	HEMBA1003908	F-HEMBA1003908	563	R-HEMBA1003908	6089
30	HEMBA1003926	F-HEMBA1003926	564	R-HEMBA1003926	6090
	HEMBA1003937	F-HEMBA1003937	565	R-HEMBA1003937	6091
	HEMBA1003939	F-HEMBA1003939	566	R-HEMBA1003939	6092
	HEMBA1003942	F-HEMBA1003942	567	R-HEMBA1003942	6093
55					

	HEMBA1003950	F-HEMBA1003950	568	R-HEMBA1003950	6094
	HEMBA1003953	F-HEMBA1003953	569	R-HEMBA1003953	6095
	HEMBA1003958	F-HEMBA1003958	570	R-HEMBA1003958	6096
5	HEMBA1003959	F-HEMBA1003959	571	R-HEMBA1003959	6097
	HEMBA1003976	F-HEMBA1003976	572	R-HEMBA1003976	6098
	HEMBA1003978	F-HEMBA1003978	573		
				R-HEMBA1003978	6099
	HEMBA1003985	F-HEMBA1003985	574 575	R-HEMBA1003985	6100
10	HEMBA1003987	F-HEMBA1003987	575	R-HEMBA1003987	6101
	HEMBA1003989	F-HEMBA1003989	576	R-HEMBA1003989	6102
	HEMBA1004000	F-HEMBA1004000	577	R-HEMBA1004000	6103
	HEMBA1004011	F-HEMBA1004011	578	R-HEMBA1004011	6104
15	HEMBA1004012	F-HEMBA1004012	579	R-HEMBA1004012	6105
	HEMBA1004015	F-HEMBA1004015	580	R-HEMBA1004015	6106
	HEMBA1004024	F-HEMBA1004024,	581	R-HEMBA1004024	6107
	HEMBA1004038	F-HEMBA1004038	582	R-HEMBA1004038	6108
20	HEMBA1004042	F-HEMBA1004042	583	R-HEMBA1004042	6109
20	HEMBA1004045	F-HEMBA1004045	584	R-HEMBA1004045	6110
	HEMBA1004048	F-HEMBA1004048	585	R-HEMBA1004048	6111
	HEMBA1004049	F-HEMBA1004049	586	R-HEMBA1004049	6112
	HEMBA1004055	F-HEMBA1004055	587	R-HEMBA1004055	6113
25	HEMBA1004056	F-HEMBA1004056	588	R-HEMBA1004056	6114
	HEMBA1004074	F-HEMBA1004074	589	R-HEMBA1004074	6115
	HEMBA1004086	F-HEMBA1004086	590	R-HEMBA1004086	6116
	HEMBA1004097	F-HEMBA1004097	591	R-HEMBA1004097	6117
30	HEMBA1004111	F-HEMBA1004111	592		
	HEMBA1004131	F-HEMBA1004131	593	R-HEMBA1004131	6118
	HEMBA1004132	F-HEMBA1004132	594	R-HEMBA1004132	6119
	HEMBA1004133	F-HEMBA1004133	595	R-HEMBA1004133	6120
	HEMBA1004138	F-HEMBA1004138	596	R-HEMBA1004138	6121
35	HEMBA1004143	F-HEMBA1004143	597	R-HEMBA1004143	6122
	HEMBA1004146	F-HEMBA1004146	598	R-HEMBA1004146	6123
	HEMBA1004150	F-HEMBA1004150	599	R-HEMBA1004150	6124
	HEMBA1004164	F-HEMBA1004164	600	R-HEMBA1004164	6125
40	HEMBA1004168	F-HEMBA1004168	601	R-HEMBA1004168	6126
	HEMBA1004199	F-HEMBA1004199	602	R-HEMBA1004199	6127
	HEMBA1004200	F-HEMBA1004200	603		
	HEMBA1004200	F-HEMBA1004200		R-HEMBA1004200	6128
45			604	R-HEMBA1004202	6129
40	HEMBA1004203 HEMBA1004207	F-HEMBA1004203	605	R-HEMBA1004203	6130
		F-HEMBA1004207	606	R-HEMBA1004207	6131
	HEMBA1004225	F-HEMBA1004225	607	R-HEMBA1004225	6132
	HEMBA1004227	F-HEMBA1004227	608	R-HEMBA1004227	6133
50	HEMBA1004238	F-HEMBA1004238	609	R-HEMBA1004238	6134
	HEMBA1004241	F-HEMBA1004241	610	R-HEMBA1004241	6135
	HEMBA1004246	F-HEMBA1004246	611	R-HEMBA1004246	6136
	HEMBA1004248	F-HEMBA1004248	612	R-HEMBA1004248	6137
55	HEMBA1004264	F-HEMBA1004264	613	R-HEMBA1004264	6138

	HEMBA1004267	F-HEMBA1004267	614 ·	R-HEMBA1004267	6139
	HEMBA1004272	F-HEMBA1004272	615	R-HEMBA1004272	6140
	HEMBA1004274	F-HEMBA1004274	616	K ILMBRIOO1212	0110
5	HEMBA1004275	F-HEMBA1004275	617	R-nnnnnnnnnnn	6141
	HEMBA1004276	F-HEMBA1004276	618	R-HEMBA1004276	6142
	HEMBA1004286	F-HEMBA1004286	619	R-HEMBA1004286	6143
	HEMBA1004289	F-HEMBA1004289	620	R-HEMBA1004289	6144
	HEMBA1004289	F-HEMBA1004295	621	R-HEMBA1004285	6145
10		F-HEMBA1004295	622	R-HEMBA1004306	6146
	HEMBA1004306		623	R-HEMBA1004312	6147
	HEMBA1004312	F-HEMBA1004312	624	R-HEMBA1004321	6148
	HEMBA1004321	F-HEMBA1004321		R-HEMBA1004323	6149
15	HEMBA1004323	F-HEMBA1004323	625		
	HEMBA1004327	F-HEMBA1004327	626	R-HEMBA1004327	6150
	HEMBA1004330	F-HEMBA1004330,	627	R-HEMBA1004330	6151
	HEMBA1004334	F-HEMBA1004334	628	R-HEMBA1004334	6152
20	HEMBA1004335	F-HEMBA1004335	629	R-HEMBA1004335	6153
	HEMBA1004341	F-HEMBA1004341	630	R-HEMBA1004341	6154
	HEMBA1004353	F-HEMBA1004353	631	R-HEMBA1004353	6155
	HEMBA1004354	F-HEMBA1004354	632	R-HEMBA1004354	6156
0.5	HEMBA1004356	F-HEMBA1004356	633	R-HEMBA1004356	6157
25	HEMBA1004366	F-HEMBA1004366	634	R-HEMBA1004366	6158
	HEMBA1004372	F-HEMBA1004372	635	R-HEMBA1004372	6159
	HEMBA1004389	F-HEMBA1004389	636	R-HEMBA1004389	6160
	HEMBA1004394	F-HEMBA1004394	637	R-HEMBA1004394	6161
30	HEMBA1004396	F-HEMBA1004396	638	R-HEMBA1004396	6162
	HEMBA1004405	F-HEMBA1004405	639	R-HEMBA1004405	6163 6164
	HEMBA1004408	F-HEMBA1004408	640	R-HEMBA1004408	6165
	HEMBA1004429	F-HEMBA1004429	641	R-HEMBA1004429	6166
35	HEMBA1004433	F-HEMBA1004433	642	R-HEMBA1004433	6167
	HEMBA1004460	F-HEMBA1004460	643	R-HEMBA1004460	6168
	HEMBA1004461	F-HEMBA1004461	644	R-HEMBA1004461 R-HEMBA1004479	6169
	HEMBA1004479	F-HEMBA1004479	645	R-HEMBA1004482	6170
40	HEMBA1004482 HEMBA1004499	F-HEMBA1004482 F-HEMBA1004499	646 647	R-HEMDATOU4402	0110
,,,	HEMBA1004502	F-HEMBA1004502	648	R-HEMBA1004502	6171
		F-HEMBA1004506	649	R-HEMBA1004506	6172
	HEMBA1004507	F-HEMBA1004507	650	R-HEMBA1004507	6173
	HEMBA1004509	F-HEMBA1004509	651	R-HEMBA1004509	6174
45	HEMBA1004534	F-HEMBA1004534	652	R-HEMBA1004534	6175
	HEMBA1004538	F-HEMBA1004538	653	R-HEMBA1004538	6176
	HEMBA1004542	F-HEMBA1004542	654	K-HEMDY1004990	0110
	HEMBA1004554	F-HEMBA1004554	655	R-HEMBA1004554	6177
50	HEMBA1004560	F-HEMBA1004560	656	R-HEMBA1004560	6178
			657	R-HEMBA1004500	6179
	HEMBA1004573	F-HEMBA1004573 F-HEMBA1004577	658	R-HEMBA1004577	6180
	HEMBA1004577	F-HEMBA1004586	659	R-HEMBA1004577	6181
55	HEMBA1004586	L-Urany I AA 4000	005	V HENDATOV-1000	0101

	HEMBA1004596	F-HEMBA1004596	660	R-nnnnnnnnnn	6182
	HEMBA1004604	F-HEMBA1004604	661		
•	HEMBA1004610	F-HEMBA1004610	662	R-HEMBA1004610	6183
5	HEMBA1004617	F-HEMBA1004617	663	R-HEMBA1004617	6184
	HEMBA1004629	F-HEMBA1004629	664	R-HEMBA1004629	6185
	HEMBA1004631	F-HEMBA1004631	665	R-HEMBA1004631	6186
	HEMBA1004632	F-HEMBA1004632	666	R-HEMBA1004632	6187
10	HEMBA1004637	F-HEMBA1004637	667	R-HEMBA1004637	6188
	HEMBA1004638	F-HEMBA1004638	668	R-HEMBA1004638	6189
	HEMBA1004666	F-HEMBA1004666	669	R-HEMBA1004666	6190
	HEMBA1004669	F-HEMBA1004669	670	R-HEMBA1004669	6191
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	HEMBA1004672	F-HEMBA1004672	672	R-HEMBA1004672	6193
	HEMBA1004693	F-HEMBA1004693,	673	R-HEMBA1004693	6194
	HEMBA1004697	F-HEMBA1004697	674	R-HEMBA1004697	6195
•	HEMBA1004705	F-HEMBA1004705	675	R-HEMBA1004705	6196
20	HEMBA1004709	F-HEMBA1004709	676	R-HEMBA1004709	6197
	HEMBA1004711	F-HEMBA1004711	677	R-HEMBA1004711	6198
	HEMBA1004725	F-HEMBA1004725	678	R-HEMBA1004725	6199
	HEMBA1004730	F-HEMBA1004730	679	R-HEMBA1004730	6200
25	HEMBA1004733	F-HEMBA1004733	680	R-HEMBA1004733	6201
	HEMBA1004734	F-HEMBA1004734	681	R-HEMBA1004734	6202
	HEMBA1004736	F-HEMBA1004736	682	R-HEMBA1004736	6203
	HEMBA1004748	F-HEMBA1004748	683	R-HEMBA1004748	6204
30	HEMBA1004751	F-HEMBA1004751	684	R-HEMBA1004751	6205
	HEMBA1004752	F-HEMBA1004752	685	R-HEMBA1004752	6206 6207
	HEMBA1004753	F-HEMBA1004753 F-HEMBA1004756	686 687	R-HEMBA1004753 R-HEMBA1004756	6208
	HEMBA1004756 HEMBA1004758	F-HEMBA1004758	688	R-HEMBA1004758	6209
35	HEMBA1004753	F-HEMBA1004763	689	R-HEMBA1004763	6210
	HEMBA1004768	F-HEMBA1004768	690	R-HEMBA1004768	6211
	HEMBA1004770	F-HEMBA1004770	691	R-HEMBA1004770	6212
	HEMBA1004771	F-HEMBA1004771	692	R-HEMBA1004771	6213
40	HEMBA1004776	F-HEMBA1004776	693	R-HEMBA1004776	6214
	HEMBA1004778	F-HEMBA1004778	694	R-HEMBA1004778	6215
	HEMBA1004795	F-HEMBA1004795	695	R-nnnnnnnnnnn	6216
	HEMBA1004803	F-HEMBA1004803	696	R-HEMBA1004803	6217
45	HEMBA1004806	F-HEMBA1004806	697	R-HEMBA1004806	6218
	HEMBA1004807	F-HEMBA1004807	698	R-HEMBA1004807	6219
	HEMBA1004816	F-HEMBA1004816	699	R-HEMBA1004816	6220
	HEMBA1004820	F-HEMBA1004820	700	R-HEMBA1004820	6221
50	HEMBA1004847	F-HEMBA1004847	701	R-HEMBA1004847	6222
50	HEMBA1004850	F-HEMBA1004850	702	R-HEMBA1004850	6223
	HEMBA1004863	F-HEMBA1004863	703	R-HEMBA1004863	6224
	HEMBA1004864	F-HEMBA1004864	704	R-HEMBA1004864	6225
	HEMBA1004865	F-HEMBA1004865	705	R-HEMBA1004865	6226
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	HEMBA1004880	F-HEMBA1004880	706	R-HEMBA1004880	6227
	HEMBA1004889	F-HEMBA1004889	707	R-HEMBA1004889	6228
	HEMBA1004900	F-HEMBA1004900	708	R-HEMBA1004900	6229
5	HEMBA1004909	F-HEMBA1004909	709	R-HEMBA1004909	6230
	HEMBA1004918	F-HEMBA1004918	710	R-HEMBA1004918	6231
	HEMBA1004923	F-HEMBA1004923	711	R-HEMBA1004923	6232
	HEMBA1004929	F-HEMBA1004929	712	R-HEMBA1004929	6233
10	HEMBA1004930	F-HEMBA1004930	713	R-HEMBA1004930	6234
	HEMBA1004933	F-HEMBA1004933	714	R-HEMBA1004933	6235
	HEMBA1004934	F-HEMBA1004934	715	R-HEMBA1004934	6236
	HEMBA1004944	F-HEMBA1004944	716	R-HEMBA1004944	6237
45	HEMBA1004954	F-HEMBA1004954	717	R-HEMBA1004954	6238
15	HEMBA1004956	F-HEMBA1004956	718	R-HEMBA1004956	6239
	HEMBA1004960	F-HEMBA1004960,	719	R-HEMBA1004960	6240
	HEMBA1004972	F-HEMBA1004972	720	R-HEMBA1004972	6241
	HEMBA1004973	F-HEMBA1004973	721	R-HEMBA1004973	6242
20	HEMBA1004977	F-HEMBA1004977	722	R-HEMBA1004977	6243
	HEMBA1004978	F-HEMBA1004978	723	R-HEMBA1004978	6244
	HEMBA1004980	F-HEMBA1004980	724	R-HEMBA1004980	6245
	HEMBA1004983	F-HEMBA1004983	725	R-HEMBA1004983	6246
25	HEMBA1004995	F-HEMBA1004995	726	R-HEMBA1004995	6247
	HEMBA1005008	F-HEMBA1005008	727	R-HEMBA1005008	6248
	HEMBA1005009	F-HEMBA1005009	728	R-HEMBA1005009	6249
	HEMBA1005019	F-HEMBA1005019	729	R-HEMBA1005019	6250
30	HEMBA1005029	F-HEMBA1005029	730	R-HEMBA1005029	6251
	HEMBA1005035	F-HEMBA1005035	731	R-HEMBA1005035	6252
	HEMBA1005039	F-HEMBA1005039	732	R-HEMBA1005039	6253
	HEMBA1005047	F-HEMBA1005047	733	R-HEMBA1005047	6254
35	HEMBA1005050	F-HEMBA1005050	734	R-HEMBA1005050	6255
	HEMBA1005062	F-HEMBA1005062	735	R-HEMBA1005062	6256
	HEMBA1005066	F-HEMBA1005066	736	R-HEMBA1005066	6257
	HEMBA1005075	F-HEMBA1005075	737	R-HEMBA1005075	6258
40	HEMBA1005079	F-HEMBA1005079	738	R-HEMBA1005079	6259
40	HEMBA1005083 HEMBA1005101	F-HEMBA1005083	739	R-HEMBA1005083	6260 6261
	HEMBA1005113	F-HEMBA1005101 F-HEMBA1005113	740	R-HEMBA1005101	6262
	HEMBA1005123	F-HEMBA1005113	741	R-HEMBA1005113	6263
	HEMBA1005123	F-HEMBA1005123	742 743	R-HEMBA1005123 R-HEMBA1005133	6264
45	HEMBA1005149	F-HEMBA1005149	743 744	R-HEMBA1005149	6265
	HEMBA1005152	F-HEMBA1005152	744 745	R-HEMBA1005152	6266
	HEMBA1005159	F-HEMBA1005152	745 746	R-HEMBA1005159	6267
	HEMBA1005185	F-HEMBA1005185	740 747	R-HEMBA1005185	6268
50	HEMBA1005201	F-HEMBA1005201	748	R-HEMBA1005201	6269
	HEMBA1005201	F-HEMBA1005201	749	R-HEMBA1005201	6270
	HEMBA1005202	F-HEMBA1005202	750	K HEMBRIVOZOZ	0210
	HEMBA1005219	F-HEMBA1005219	751	R-HEMBA1005219	6271
55	11L8IDA 1003219	I. Ilrum.(1002512	101	it immittooos13	0211

	HEMBA1005223	F-HEMBA1005223	752	R-HEMBA1005223	6272
	HEMBA1005232	F-HEMBA1005232	753	R-HEMBA1005232	6273
	HEMBA1005241	F-HEMBA1005241	754	R-HEMBA1005241	6274
5	HEMBA1005244	F-HEMBA1005244	755	R-HEMBA1005244	6275
	HEMBA1005251	F-HEMBA1005251	756	R-HEMBA1005251	6276
	HEMBA1005252	F-HEMBA1005252	757	R-HEMBA1005252	6277
	HEMBA1005274	F-HEMBA1005274	758	R-HEMBA1005274	6278
	HEMBA1005275	F-HEMBA1005275	759	R-HEMBA1005275	6279
10	HEMBA1005293	F-HEMBA1005293	760	R-HEMBA1005293	6280
	HEMBA1005296	F-HEMBA1005296	761	R-HEMBA1005296	6281
	HEMBA1005304	F-HEMBA1005304	762	R-HEMBA1005304	6282
	HEMBA1005304	F-HEMBA1005311	763	R-HEMBA1005311	6283
15					6284
	HEMBA1005314	F-HEMBA1005314	764	R-HEMBA1005314	6285
	HEMBA1005315	F-HEMBA1005315,	765	R-HEMBA1005315	
	HEMBA1005318	F-HEMBA1005318	766	R-HEMBA1005318	6286
20	HEMBA1005331	F-HEMBA1005331	767	R-HEMBA1005331	6287
20	HEMBA1005338	F-HEMBA1005338	768	n imim. 1005050	2000
	HEMBA1005353	F-HEMBA1005353	769	R-HEMBA1005353	6288
	HEMBA1005359	F-HEMBA1005359	770	R-HEMBA1005359	6289
	HEMBA1005367	F-HEMBA1005367	771	R-HEMBA1005367	6290
25	HEMBA1005372	F-HEMBA1005372	772	R-HEMBA1005372	6291
	HEMBA1005374	F-HEMBA1005374	773	R-HEMBA1005374	6292
	HEMBA1005382	F-HEMBA1005382	774		
	HEMBA1005389	F-HEMBA1005389	775	R-HEMBA1005389	6293
30	HEMBA1005394	F-HEMBA1005394	776	R-HEMBA1005394	6294
	HEMBA1005403	F-HEMBA1005403	777	R-HEMBA1005403	6295
	HEMBA1005408	F-HEMBA1005408	778	R-HEMBA1005408	6296
	HEMBA1005410	F-HEMBA1005410	779	R-HEMBA1005410	6297
25	HEMBA1005411	F-HEMBA1005411	780	R-HEMBA1005411	6298
35	HEMBA1005423	F-HEMBA1005423	781	R-HEMBA1005423	6299
	HEMBA1005426	F-HEMBA1005426	782	R-HEMBA1005426	6300
	HEMBA1005443	F-HEMBA1005443	783	R-HEMBA1005443	6301
	HEMBA1005447	F-HEMBA1005447	784	R-HEMBA1005447	6302
40	HEMBA1005468	F-HEMBA1005468	785	R-HEMBA1005468	6303
	HEMBA1005469	F-HEMBA1005469	786	R-HEMBA1005469	6304
	HEMBA1005472	F-HEMBA1005472	787	R-HEMBA1005472	6305
	HEMBA1005474	F-HEMBA1005474	788		
45	HEMBA1005475	F-HEMBA1005475	789	R-HEMBA1005475	6306
	HEMBA1005497	F-HEMBA1005497	790	R-HEMBA1005497	6307
	HEMBA1005500	F-HEMBA1005500	791	R-HEMBA1005500	6308
	HEMBA1005506	F-HEMBA1005506	792	R-HEMBA1005506	6309
	HEMBA1005508	F-HEMBA1005508	793	R-HEMBA1005508	6310
50	HEMBA1005511	F-HEMBA1005511	794	R-HEMBA1005511	6311
	HEMBA1005513	F-HEMBA1005513	795		
	HEMBA1005517	F-HEMBA1005517	796	R-HEMBA1005517	6312
	HEMBA1005518	F-HEMBA1005518	797	R-HEMBA1005518	6313
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	HEMBA1005520	F-HEMBA1005520	798	R-HEMBA1005520	6314
	HEMBA1005526	F-HEMBA1005526	799	R-HEMBA1005526	6315
	HEMBA1005528	F-HEMBA1005528	800	R-HEMBA1005528	6316
5	HEMBA1005530	F-HEMBA1005530	801	R-HEMBA1005530	6317
	HEMBA1005548	F-HEMBA1005548	802	R-HEMBA1005548	6318
	HEMBA1005552	F-HEMBA1005552	803	R-HEMBA1005552	6319
	HEMBA1005558	F-HEMBA1005558	804	R-HEMBA1005558	6320
10	HEMBA1005568	F-HEMBA1005568	805	R-HEMBA1005568	6321
	HEMBA1005570	F-HEMBA1005570	806	R-HEMBA1005570	6322
	HEMBA1005576	F-HEMBA1005576	807	R-HEMBA1005576	6323
	HEMBA1005577	F-HEMBA1005577	808	R-HEMBA1005577	6324
15	HEMBA1005581	F-HEMBA1005581	809	R-HEMBA1005581	6325
75	HEMBA1005582	F-HEMBA1005582	810	R-HEMBA1005582	6326
	HEMBA1005583	F-HEMBA1005583,	811	R-HEMBA1005583	6327
	HEMBA1005588	F-HEMBA1005588	812	R-HEMBA1005588	6328
	HEMBA1005593	F-HEMBA1005593	813	R-HEMBA1005593	6329
20	HEMBA1005595	F-HEMBA1005595	814	R-HEMBA1005595	6330
	HEMBA1005606	F-HEMBA1005606	815	R-HEMBA1005606	6331
	HEMBA1005609	F-HEMBA1005609	816	R-HEMBA1005609	6332
	HEMBA1005616	F-HEMBA1005616	817	R-HEMBA1005616	6333
25	HEMBA1005621	F-HEMBA1005621	818	R-HEMBA1005621	6334
	HEMBA1005627	F-HEMBA1005627	819	R-HEMBA1005627	6335
	HEMBA1005631	F-HEMBA1005631	820	R-HEMBA1005631	6336
	HEMBA1005632	F-HEMBA1005632	821	R-HEMBA1005632	6337
30	HEMBA1005634	F-HEMBA1005634	822	R-HEMBA1005634	6338
	HEMBA1005666	F-HEMBA1005666	823	R-HEMBA1005666	6339
		F-HEMBA1005670	824	R-HEMBA1005670	6340
	HEMBA1005679	F-HEMBA1005679	825	R-HEMBA1005679	6341 6342
35	HEMBA1005680 HEMBA1005685	F-HEMBA1005680 F-HEMBA1005685	826 827	R-HEMBA1005680 R-HEMBA1005685	6343
	HEMBA1005699	F-HEMBA1005699	828	R-HEMBA1005699	6344
	HEMBA1005705	F-HEMBA1005705	829	R-HEMBA1005705	6345
	HEMBA1005717	F-HEMBA1005717	830	R-HEMBA1005717	6346
40	HEMBA1005732	F-HEMBA1005732	831	R-HEMBA1005732	6347
	HEMBA1005737	F-HEMBA1005737	832	R-HEMBA1005737	6348
	HEMBA1005746	F-HEMBA1005746	833	R-nnnnnnnnnnn	6349
	HEMBA1005755	F-HEMBA1005755	834	R-HEMBA1005755	6350
45	HEMBA1005765	F-HEMBA1005765	835	R-HEMBA1005765	6351
45	HEMBA1005780	F-HEMBA1005780	836	R-HEMBA1005780	6352
	HEMBA1005813	F-HEMBA1005813	837	R-HEMBA1005813	6353
	HEMBA1005815	F-HEMBA1005815	838	R-HEMBA1005815	6354
	HEMBA1005822	F-HEMBA1005822	839	R-HEMBA1005822	6355
50	HEMBA1005829	F-HEMBA1005829	840	R-HEMBA1005829	6356
	HEMBA1005834	F-HEMBA1005834	841	R-HEMBA1005834	6357
	HEMBA1005852	F-HEMBA1005852	842	R-HEMBA1005852	6358
	HEMBA1005853	F-HEMBA1005853	843	R-HEMBA1005853	6359
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	HEMBA1005884	F-HEMBA1005884	844	R-HEMBA1005884	6360
	HEMBA1005891	F-HEMBA1005891	845	R-HEMBA1005891	6361
	HEMBA1005894	F-HEMBA1005894	846	R-HEMBA1005894	6362
5	HEMBA1005909	F-HEMBA1005909	847	R-HEMBA1005909	6363
	HEMBA1005911	F-HEMBA1005911	848	R-HEMBA1005911	6364
	HEMBA1005921	F-HEMBA1005921	849	R-HEMBA1005921	6365
	HEMBA1005931	F-HEMBA1005931	850	R-HEMBA1005931	6366
10	HEMBA1005934	F-HEMBA1005934	851	R-HEMBA1005934	6367
	HEMBA1005962	F-HEMBA1005962	852	R-HEMBA1005962	6368
	HEMBA1005963	F-HEMBA1005963	853	R-HEMBA1005963	6369
	HEMBA1005990	F-HEMBA1005990	854	R-HEMBA1005990	6370
15	HEMBA1005991	F-HEMBA1005991	855	R-HEMBA1005991	6371
13	HEMBA1005999	F-HEMBA1005999	856	R-HEMBA1005999	6372
	HEMBA1006002	F-HEMBA1006002,	857	R-HEMBA1006002	6373
	HEMBA1006005	F-HEMBA1006005	858	R-HEMBA1006005	6374
	HEMBA1006031	F-HEMBA1006031	859	R-nnnnnnnnnnn	6375
20	HEMBA1006035	F-HEMBA1006035	860	R-HEMBA1006035	6376
	HEMBA1006036	F-HEMBA1006036	861	R-HEMBA1006036	6377
	HEMBA1006042	F-HEMBA1006042	862	R-HEMBA1006042	6378
	HEMBA1006067	F-HEMBA1006067	863	R-nnnnnnnnnnn	6379
25	HEMBA1006081	F-HEMBA1006081	864	R-HEMBA1006081	6380
	HEMBA1006090	F-HEMBA1006090	865	R-HEMBA1006090	6381 6382
	HEMBA1006091	F-HEMBA1006091	866	R-HEMBA1006091 R-HEMBA1006100	6383
	HEMBA1006100 HEMBA1006108	F-HEMBA1006100 F-HEMBA1006108	867 868	R-HEMBA1006108	6384
30	HEMBA1006121	F-HEMBA1006121	869	R-HEMBA1006121	6385
	HEMBA1006124	F-HEMBA1006124	870	R-HEMBA1006124	6386
	HEMBA1006130	F-HEMBA1006130	871	R-HEMBA1006130	6387
	HEMBA1006138	F-HEMBA1006138	872	R-nnnnnnnnnn	6388
35	HEMBA1006142	F-HEMBA1006142	873	R-HEMBA1006142	6389
	HEMBA1006155	F-HEMBA1006155	874	R-HEMBA1006155	6390
	HEMBA1006158	F-HEMBA1006158	875	R-HEMBA1006158	6391
	HEMBA1006173	F-HEMBA1006173	876	R-HEMBA1006173	6392
40	HEMBA1006182	F-HEMBA1006182	877	R-HEMBA1006182	6393
	HEMBA1006198	F-HEMBA1006198	878	R-HEMBA1006198	6394
	HEMBA1006235	F-HEMBA1006235	879	R-HEMBA1006235	6395
	HEMBA1006248	F-HEMBA1006248	880	R-HEMBA1006248	6396
45	HEMBA1006252	F-HEMBA1006252	881	R-HEMBA1006252	6397
	HEMBA1006253	F-HEMBA1006253	882	R-HEMBA1006253	6398
	HEMBA1006259	F-HEMBA1006259	883	R-HEMBA1006259	6399
	HEMBA1006268	F-HEMBA1006268	884	R-HEMBA1006268	6400
50	HEMBA1006272	F-HEMBA1006272	885	R-HEMBA1006272	6401
	HEMBA1006278	F-HEMBA1006278	886	R-nnnnnnnnnnnn	6402
	HEMBA1006283	F-HEMBA1006283	887 888	R-HEMBA1006283 R-HEMBA1006284	6403 6404
	HEMBA1006284	F-HEMBA1006284 F-HEMBA1006291	889	R-HEMBA1006291	6405
55	HEMBA1006291	r-nembaluu0291	003	K-IIIMDA1000291	0403
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	HEMBA1006293	F-HEMBA1006293	890	R-HEMBA1006293	· 6406
	HEMBA1006309	F-HEMBA1006309	891	R-HEMBA1006309	6407
	HEMBA1006310	F-HEMBA1006310	892	R-HEMBA1006310	6408
5	HEMBA1006328	F-HEMBA1006328	893	R-HEMBA1006328	6409
	HEMBA1006334	F-HEMBA1006334	894	R-HEMBA1006334	6410
	HEMBA1006344	F-HEMBA1006344	895	R-HEMBA1006344	6411
	HEMBA1006347	F-HEMBA1006347	896	R-HEMBA1006347	6412
10	HEMBA1006349	F-HEMBA1006349	897	R-HEMBA1006349	6413
	HEMBA1006359	F-HEMBA1006359	898	R-HEMBA1006359	6414
	HEMBA1006364	F-HEMBA1006364	899	R-HEMBA1006364	6415
	HEMBA1006377	F-HEMBA1006377	900	R-HEMBA1006377	6416
	HEMBA1006380	F-HEMBA1006380	901	R-HEMBA1006380	6417
15	HEMBA1006381	F-HEMBA1006381	902	R-HEMBA1006381	6418
	HEMBA1006398	F-HEMBA1006398,	903	R-HEMBA1006398	6419
	HEMBA1006416	F-HEMBA1006416	904	R-HEMBA1006416	6420
	HEMBA1006419	F-HEMBA1006419	905	R-HEMBA1006419	6421
20	HEMBA1006421	F-HEMBA1006421	906	R-HEMBA1006421	6422
	HEMBA1006424	F-HEMBA1006424	907	R-HEMBA1006424	6423
	HEMBA1006426	F-HEMBA1006426	908	R-HEMBA1006426	6424
	HEMBA1006438	F-HEMBA1006438	909	R-HEMBA1006438	6425
25	HEMBA1006445	F-HEMBA1006445	910	R-HEMBA1006445	6426
	HEMBA1006446	F-HEMBA1006446	911	R-HEMBA1006446	6427
	HEMBA1006461	F-HEMBA1006461	912	R-HEMBA1006461	6428
	HEMBA1006467	F-HEMBA1006467	913	R-HEMBA1006467	6429
20	HEMBA1006471	F-HEMBA1006471	914	R-HEMBA1006471	6430
30	HEMBA1006474	F-HEMBA1006474	915	R-HEMBA1006474	6431
	HEMBA1006483	F-HEMBA1006483	916	R-HEMBA1006483	6432
	HEMBA1006485	F-HEMBA1006485	917	R-HEMBA1006485	6433
	HEMBA1006486	F-HEMBA1006486	918	R-HEMBA1006486	6434
35	HEMBA1006489	F-HEMBA1006489	919	R-HEMBA1006489	6435
	HEMBA1006492	F-HEMBA1006492	920	R-HEMBA1006492	6436
	HEMBA1006494	F-HEMBA1006494	921	R-HEMBA1006494	6437
	HEMBA1006497	F-HEMBA1006497	922	R-HEMBA1006497	6438
40	HEMBA1006502	F-HEMBA1006502	923	R-HEMBA1006502	6439
	HEMBA1006507	F-HEMBA1006507	924	R-HEMBA1006507	6440
	HEMBA1006521	F-HEMBA1006521	925	R-HEMBA1006521	6441
	HEMBA1006530	F-HEMBA1006530	926	R-HEMBA1006530	6442
45	HEMBA1006535	F-HEMBA1006535	927	R-HEMBA1006535	6443
40	HEMBA1006540	F-HEMBA1006540	928	R-HEMBA1006540	6 444
	HEMBA1006546	F-HEMBA1006546	929	R-HEMBA1006546	6445
	HEMBA1006559	F-HEMBA1006559	930	R-HEMBA1006559	6446
	HEMBA1006562	F-HEMBA1006562	931	R-HEMBA1006562	6447
50	HEMBA1006566	F-HEMBA1006566	932	R-HEMBA1006566	6448
	HEMBA1006569	F-HEMBA1006569	933	R-HEMBA1006569	6449
	HEMBA1006579	F-HEMBA1006579	934	R-HEMBA1006579	6450
	HEMBA1006583	F-HEMBA1006583	935	R-HEMBA1006583	6451

	HEMBA1006595	F-HEMBA1006595	936	R-HEMBA1006595	6452
	HEMBA1006597	F-HEMBA1006597	937	R-HEMBA1006597	6453
	HEMBA1006612	F-HEMBA1006612	938	R-HEMBA1006612	6454
5	HEMBA1006617	F-HEMBA1006617	939	R-nnnnnnnnnnn	6455
	HEMBA1006624	F-HEMBA1006624	940	R-HEMBA1006624	6456
	HEMBA1006631	F-HEMBA1006631	941	R-HEMBA1006631	6457
	HEMBA1006635	F-HEMBA1006635	942	R-HEMBA1006635	6458
40	HEMBA1006639	F-HEMBA1006639	943	R-HEMBA1006639	6459
10	HEMBA1006643	F-HEMBA1006643	944	R-HEMBA1006643	6460
	HEMBA1006648	F-HEMBA1006648	945	R-HEMBA1006648	6461
	HEMBA1006652	F-HEMBA1006652	946	R-HEMBA1006652	6462
	HEMBA1006653	F-HEMBA1006653	947	R-HEMBA1006653	6463
15	HEMBA1006659	F-HEMBA1006659	948	It insubitioned	0100
	HEMBA1006665	F-HEMBA1006665	949	R-HEMBA1006665	6464
	HEMBA1006674	F-HEMBA1006674	950	R-HEMBA1006674	6465
	HEMBA1006676	F-HEMBA1006676	951	R-HEMBA1006676	6466
20	HEMBA1006682	F-HEMBA1006682	952	R-HEMBA1006682	6467
	HEMBA1006695	F-HEMBA1006695	953	R-HEMBA1006695	6468
	HEMBA1006696	F-HEMBA1006696	954	R-HEMBA1006696	6469
	HEMBA1006708	F-HEMBA1006708	955	R-HEMBA1006708	6470
25	HEMBA1006709	F-HEMBA1006709	956	R-HEMBA1006709	6471
23	HEMBA1006717	F-HEMBA1006717	957	R-HEMBA1006717	6472
	HEMBA1006737	F-HEMBA1006737	958	R-HEMBA1006737	6473
	HEMBA1006744	F-HEMBA1006744	959	R-HEMBA1006744	6474
	HEMBA1006754	F-HEMBA1006754	960	R-HEMBA1006754	6475
30	HEMBA1006758	F-HEMBA1006758	961	R-HEMBA1006758	6476
	HEMBA1006767	F-HEMBA1006767	962	R-HEMBA1006767	6477
	HEMBA1006779	F-HEMBA1006779	963	R-HEMBA1006779	6478
	HEMBA1006780	F-HEMBA1006780	964	R-HEMBA1006780	6479
35	HEMBA1006789	F-HEMBA1006789	965	R-HEMBA1006789	6480
	HEMBA1006795	F-HEMBA1006795	966	R-HEMBA1006795	6481
	HEMBA1006796	F-HEMBA1006796	967	R-HEMBA1006796	6482
	HEMBA1006807	F-HEMBA1006807	968	R-HEMBA1006807	6483
40	HEMBA1006821	F-HEMBA1006821	969	R-HEMBA1006821	6484
	HEMBA1006824	F-HEMBA1006824	970	R-HEMBA1006824	6485
	HEMBA1006832	F-HEMBA1006832	971	R-HEMBA1006832	6486
	HEMBA1006849	F-HEMBA1006849	972	R-HEMBA1006849	6487
46	HEMBA1006865	F-HEMBA1006865	973	R-HEMBA1006865	6488
45	HEMBA1006877	F-HEMBA1006877	974	R-nnnnnnnnnnn	6489
	HEMBA1006885	F-HEMBA1006885	975	R-HEMBA1006885	6490
	HEMBA1006900	F-HEMBA1006900	976	R-HEMBA1006900	6491
	HEMBA1006914	F-HEMBA1006914	977		
50	HEMBA1006921	F-HEMBA1006921	978	R-HEMBA1006921	6492
	HEMBA1006926	F-HEMBA1006926	979	R-HEMBA1006926	6493
	HEMBA1006929	F-HEMBA1006929	980	R-HEMBA1006929	6494
	HEMBA1006936	F-HEMBA1006936	981	R-HEMBA1006936	6495

	HEMBA1006938	F-HEMBA1006938	982	R-HEMBA1006938	6496
	HEMBA1006941	F-HEMBA1006941	983	R-HEMBA1006941	6497
	HEMBA1006949	F-HEMBA1006949	984	R-HEMBA1006949	6498
5	HEMBA1006973	F-HEMBA1006973	985	R-HEMBA1006973	6499
	HEMBA1006976	F-HEMBA1006976	986	R-HEMBA1006976	6500
	HEMBA1006993	F-HEMBA1006993	987	R-HEMBA1006993	6501
	HEMBA1006996	F-HEMBA1006996	988	R-HEMBA1006996	6502
10	HEMBA1007002	F-HEMBA1007002	989	R-HEMBA1007002	6503
	HEMBA1007017	F-HEMBA1007017	990	R-HEMBA1007017	6504
	HEMBA1007018	F-HEMBA1007018	991	R-HEMBA1007018	6505
	HEMBA1007045	F-HEMBA1007045	992	R-HEMBA1007045	6506
4-	HEMBA1007051	F-HEMBA1007051	993	R-HEMBA1007051	6507
15	HEMBA1007052	F-HEMBA1007052	994	R-HEMBA1007052	6508
	HEMBA1007062	F-HEMBA1007062,	995	R-HEMBA1007062	6509
	HEMBA1007066	F-HEMBA1007066	996	R-HEMBA1007066	6510
	HEMBA1007073	F-HEMBA1007073	997	R-HEMBA1007073	6511
20	HEMBA1007078	F-HEMBA1007078	998	R-HEMBA1007078	6512
	HEMBA1007080	F-HEMBA1007080	999		
	HEMBA1007085	F-HEMBA1007085	1000	R-HEMBA1007085	6513
	HEMBA1007087	F-HEMBA1007087	1001	R-HEMBA1007087	6514
25	HEMBA1007112	F-HEMBA1007112	1002	R-HEMBA1007112	6515
	HEMBA1007113	F-HEMBA1007113	1003	R-HEMBA1007113	6516
	HEMBA1007121	F-HEMBA1007121	1004		
	HEMBA1007129	F-HEMBA1007129	1005	R-HEMBA1007129	6517
30	HEMBA1007147	F-HEMBA1007147	1006	R-HEMBA1007147	6518
30	HEMBA1007149	F-HEMBA1007149	1007	R-HEMBA1007149	6519
	HEMBA1007151	F-HEMBA1007151	1008	R-HEMBA1007151	6520
	HEMBA1007174	F-HEMBA1007174	1009	R-nnnnnnnnnnn	6521
	HEMBA1007178	F-HEMBA1007178	1010	R-HEMBA1007178	6522
35	HEMBA1007194	F-HEMBA1007194	1011	R-HEMBA1007194	6523
	HEMBA1007203	F-HEMBA1007203	1012	R-HEMBA1007203	6524
	HEMBA1007206	F-HEMBA1007206	1013	R-HEMBA1007206	6525
	HEMBA1007224	F-HEMBA1007224	1014	R-HEMBA1007224	6526
40	HEMBA1007243	F-HEMBA1007243	1015		
	HEMBA1007251	F-HEMBA1007251	1016	R-HEMBA1007251	6527
	HEMBA1007256	F-HEMBA1007256	1017	R-HEMBA1007256	6528
	HEMBA1007267	F-HEMBA1007267	1018	R-HEMBA1007267	6529
45	HEMBA1007273	F-HEMBA1007273	1019	R-HEMBA1007273	6530
	HEMBA1007279	F-HEMBA1007279	1020	R-HEMBA1007279	6531
	HEMBA1007281	F-HEMBA1007281	1021	R-HEMBA1007281	6532
	HEMBA1007288	F-HEMBA1007288	1022	R-HEMBA1007288	6533
	HEMBA1007300	F-HEMBA1007300	1023	R-HEMBA1007300	6534
50	HEMBA1007301	F-HEMBA1007301	1024	R-HEMBA1007301	6535
	HEMBA1007319	F-HEMBA1007319	1025	R-HEMBA1007319	6536
	HEMBA1007320	F-HEMBA1007320	1026	R-HEMBA1007320	6537
	HEMBA1007322	F-HEMBA1007322	1027	R-HEMBA1007322	6538

	HEMBA1007327	F-HEMBA1007327	1028	R-HEMBA1007327	6539
	HEMBA1007341	F-HEMBA1007341	1029	R-HEMBA1007341	6540
	HEMBA1007342	F-HEMBA1007342	1030	R-HEMBA1007342	6541
5	HEMBA1007347	F-HEMBA1007347	1031	R-HEMBA1007347	6542
	HEMBB1000005	F-HEMBB1000005	1032	R-HEMBB1000005	6543
	HEMBB1000008	F-HEMBB1000008	1033	R-HEMBB1000008	6544
	HEMBB1000018	F-HEMBB1000018	1034	R-HEMBB1000018	6545
10	HEMBB1000024	F-HEMBB1000024	1035	R-HEMBB1000024	6546
	HEMBB1000025	F-HEMBB1000025	1036	R-HEMBB1000025	6547
	HEMBB1000030	F-HEMBB1000030	1037	R-HEMBB1000030	6548
	HEMBB1000036	F-HEMBB1000036	1038	R-HEMBB1000036	6549
	HEMBB1000037	F-HEMBB1000037	1039	R-HEMBB1000037	6550
15	HEMBB1000039	F-HEMBB1000039	1040	R-HEMBB1000039	6551
	HEMBB1000044	F-HEMBB1000044,	1041	R-HEMBB1000044	6552
	HEMBB1000048	F-HEMBB1000048	1042	R-HEMBB1000048	6553
	HEMBB1000050	F-HEMBB1000050	1043	R-HEMBB1000050	6554
20	HEMBB1000054	F-HEMBB1000054	1044	R-HEMBB1000054	6555
	HEMBB1000055	F-HEMBB1000055	1045	R-HEMBB1000055	6556
	HEMBB1000059	F-HEMBB1000059	1046	R-HEMBB1000059	6557
	HEMBB1000083	F-HEMBB1000083	1047	R-HEMBB1000083	6558
25	HEMBB1000089	F-HEMBB1000089	1048	R-HEMBB1000089	6559
	HEMBB1000099	F-HEMBB1000099	1049	R-HEMBB1000099	6560
	HEMBB1000103	F-HEMBB1000103	1050	R-HEMBB1000103	6561
	HEMBB1000113	F-HEMBB1000113	1051	R-HEMBB1000113	6562
	HEMBB1000119	F-HEMBB1000119	1052	R-HEMBB1000119	6563
30	HEMBB1000136	F-HEMBB1000136	1053	R-HEMBB1000136	6564
	HEMBB1000141	F-HEMBB1000141	1054	R-HEMBB1000141	6565
	HEMBB1000144	F-HEMBB1000144	1055	R-HEMBB1000144	6566
	HEMBB1000173	F-HEMBB1000173	1056	R-HEMBB1000173	6567
35	HEMBB1000175	F-HEMBB1000175	1057	R-HEMBB1000175	6568
	HEMBB1000198	F-HEMBB1000198	1058	R-HEMBB1000198	6569
	HEMBB1000215	F-HEMBB1000215	1059	R-HEMBB1000215	6570
	HEMBB1000217	F-HEMBB1000217	1060	R-HEMBB1000217	6571
40	HEMBB1000218	F-HEMBB1000218	1061	R-HEMBB1000218	6572
	HEMBB1000226	F-HEMBB1000226	1062	R-HEMBB1000226	6573
	HEMBB1000240	F-HEMBB1000240	1063	R-HEMBB1000240	6574
	HEMBB1000244	F-HEMBB1000244	1064	R-HEMBB1000244	6575
4-	HEMBB1000250	F-HEMBB1000250	1065	R-HEMBB1000250	6576
45	HEMBB1000258	F-HEMBB1000258	1066	R-HEMBB1000258	6577
	HEMBB1000264	F-HEMBB1000264	1067	R-HEMBB1000264	6578
	HEMBB1000266	F-HEMBB1000266	1068	R-HEMBB1000266	6579
	HEMBB1000272	F-HEMBB1000272	1069	R-HEMBB1000272	6580
50	HEMBB1000274	F-HEMBB1000274	1070	R-HEMBB1000274	6581
	HEMBB1000284	F-HEMBB1000284	1071	R-HEMBB1000284	6582
	HEMBB1000307	F-HEMBB1000307	1072	R-HEMBB1000307	6583
	HEMBB1000312	F-HEMBB1000312	1073	R-HEMBB1000312	6584

	HEMBB1000317	F-HEMBB1000317	1074	R-HEMBB1000317	6585
	HEMBB1000318	F-HEMBB1000318	1075	R-HEMBB1000318	6586
	HEMBB1000335	F-HEMBB1000335	1076	R-HEMBB1000335	6587
5	HEMBB1000336	F-HEMBB1000336	1077	R-HEMBB1000336	6588
	HEMBB1000337	F-HEMBB1000337	1078	R-HEMBB1000337	6589
	HEMBB1000338	F-HEMBB1000338	1079	R-HEMBB1000338	6590
	HEMBB1000339	F-HEMBB1000339	1080	R-HEMBB1000339	6591
10	HEMBB1000341	F-HEMBB1000341	1081	R-HEMBB1000341	6592
,,,	HEMBB1000343	F-HEMBB1000343	1082	R-HEMBB1000343	6593
	HEMBB1000354	F-HEMBB1000354	1083	R-HEMBB1000354	6594
	HEMBB1000369	F-HEMBB1000369	1084	R-HEMBB1000369	6595
	HEMBB1000374	F-HEMBB1000374	1085	R-HEMBB1000374	6596
15	HEMBB1000376	F-HEMBB1000376	1086	R-HEMBB1000376	6597
	HEMBB1000391	F-HEMBB1000391,	1087	R-HEMBB1000391	6598
	HEMBB1000399	F-HEMBB1000399	1088	R-HEMBB1000399	6599
	HEMBB1000402	F-HEMBB1000402	1089	R-HEMBB1000402	6600
20	HEMBB1000404	F-HEMBB1000404	1090	R-HEMBB1000404	6601
	HEMBB1000420	F-HEMBB1000420	1091	R-HEMBB1000420	6602
	HEMBB1000434	F-HEMBB1000434	1092	R-HEMBB1000434	6603
	HEMBB1000438	F-HEMBB1000438	1093	R-HEMBB1000438	6604
25	HEMBB1000441	F-HEMBB1000441	1094	R-HEMBB1000441	6605
	HEMBB1000449	F-HEMBB1000449	1095	R-HEMBB1000449	6606
	HEMBB1000455	F-HEMBB1000455	1096	R-HEMBB1000455	6607
	HEMBB1000472	F-HEMBB1000472	1097	R-HEMBB1000472	6608
30	HEMBB1000480	F-HEMBB1000480	1098	R-HEMBB1000480	6609
	HEMBB1000487	F-HEMBB1000487	1099	R-HEMBB1000487	6610
	HEMBB1000490	F-HEMBB1000490	1100	R-HEMBB1000490	6611
	HEMBB1000491	F-HEMBB1000491	1101	R-HEMBB1000491	6612
	HEMBB1000493	F-HEMBB1000493	1102	R-HEMBB1000493	6613
35	HEMBB1000510	F-HEMBB1000510	1103	R-HEMBB1000510	6614
	HEMBB1000518	F-HEMBB1000518	1104	R-HEMBB1000518	6615
	HEMBB1000523	F-HEMBB1000523	1105	R-HEMBB1000523	6616
	HEMBB1000530	F-HEMBB1000530	1106	R-HEMBB1000530	6617
40	HEMBB1000550	F-HEMBB1000550	1107	R-HEMBB1000550	6618
	HEMBB1000554	F-HEMBB1000554	1108	R-HEMBB1000554	6619
	HEMBB1000556	F-HEMBB1000556	1109	R-HEMBB1000556	6620
	HEMBB1000564	F-HEMBB1000564	1110	R-HEMBB1000564	6621
45	HEMBB1000573	F-HEMBB1000573	1111	R-HEMBB1000573	6622
	HEMBB1000575	F-HEMBB1000575	1112	R-HEMBB1000575	6623
	HEMBB1000586	F-HEMBB1000586	1113	R-HEMBB1000586	6624
	HEMBB1000589	F-HEMBB1000589	1114	R-HEMBB1000589	6625
50	HEMBB1000591	F-HEMBB1000591	1115	R-HEMBB1000591	6626
	HEMBB1000592	F-HEMBB1000592	1116	R-HEMBB1000592	6627
	HEMBB1000593	F-HEMBB1000593	1117	D_UD)(DD1000500	6600
	HEMBB1000598	F-HEMBB1000598	1118	R-HEMBB1000598	6628
	HEMBB1000623	F-HEMBB1000623	1119	R-HEMBB1000623	6629

	HEMBB1000630	F-HEMBB1000630	1120	R-HEMBB1000630	6630
	HEMBB1000631	F-HEMBB1000631	1121	R-HEMBB1000631	6631
	HEMBB1000632	F-HEMBB1000632	1122	R-HEMBB1000632	6632
5	HEMBB1000637	F-HEMBB1000637	1123	R-HEMBB1000637	6633
	HEMBB1000638	F-HEMBB1000638	1124	R-HEMBB1000638	6634
	HEMBB1000643	F-HEMBB1000643	1125	R-HEMBB1000643	6635
	HEMBB1000649	F-HEMBB1000649	1126	R-HEMBB1000649	6636
10	HEMBB1000652	F-HEMBB1000652	1127	R-HEMBB1000652	6637
	HEMBB1000665	F-HEMBB1000665	1128	R-HEMBB1000665	6638
	HEMBB1000671	F-HEMBB1000671	1129	R-HEMBB1000671	6639
	HEMBB1000673	F-HEMBB1000673	1130	R-HEMBB1000673	6640
15	HEMBB1000684	F-HEMBB1000684	1131	R-HEMBB1000684	6641
,0	HEMBB1000693	F-HEMBB1000693	1132	R-nnnnnnnnnn	6642
	HEMBB1000705	F-HEMBB1000705	1133	R-HEMBB1000705	6643
	HEMBB1000706	F-HEMBB1000706	1134	R-HEMBB1000706	6644
	HEMBB1000709	F-HEMBB1000709	1135	R-HEMBB1000709	6645
20	HEMBB1000725	F-HEMBB1000725	1136	R-HEMBB1000725	6646
	HEMBB1000726	F-HEMBB1000726	1137	R-HEMBB1000726	6647
	HEMBB1000738	F-HEMBB1000738	1138	R-HEMBB1000738	6648
	HEMBB1000749	F-HEMBB1000749	1139	R-HEMBB1000749	6649
25	HEMBB1000763	F-HEMBB1000763	1140	R-HEMBB1000763	6650
	HEMBB1000770	F-HEMBB1000770	1141	R-HEMBB1000770	6651
	HEMBB1000774	F-HEMBB1000774	1142		
	HEMBB1000781	F-HEMBB1000781	1143	R-HEMBB1000781	6652
30	HEMBB1000789	F-HEMBB1000789	1144	R-HEMBB1000789	6653
	HEMBB1000790	F-HEMBB1000790	1145	R-HEMBB1000790	6654
	HEMBB1000794	F-HEMBB1000794	1146	R-HEMBB1000794	6655
	HEMBB1000807	F-HEMBB1000807	1147	R-HEMBB1000807	6656
35	HEMBB1000810	F-HEMBB1000810	1148	R-HEMBB1000810	6657
	HEMBB1000821	F-HEMBB1000821	1149	R-HEMBB1000821	6658
•	HEMBB1000822	F-HEMBB1000822	1150	R-HEMBB1000822	6659
	HEMBB1000826	F-HEMBB1000826	1151	R-HEMBB1000826	6660
40	HEMBB1000827	F-HEMBB1000827	1152	R-HEMBB1000827	6661
	HEMBB1000831	F-HEMBB1000831	1153	R-HEMBB1000831	6662
	HEMBB1000835	F-HEMBB1000835	1154	R-HEMBB1000835	6663
	HEMBB1000840	F-HEMBB1000840	1155	R-HEMBB1000840	6664
45	HEMBB1000848	F-HEMBB1000848	1156	R-HEMBB1000848	6665
4 5	HEMBB1000852	F-HEMBB1000852	1157	R-HEMBB1000852	6666
	HEMBB1000870	F-HEMBB1000870	1158	R-HEMBB1000870	6667
	HEMBB1000876	F-HEMBB1000876	1159	R-HEMBB1000876	6668
	HEMBB1000883	F-HEMBB1000883	1160	R-HEMBB1000883	6669
50	HEMBB1000887	F-HEMBB1000887	1161	R-HEMBB1000887	6670
	HEMBB1000888	F-HEMBB1000888	1162	R-HEMBB1000888	6671
	HEMBB1000890	F-HEMBB1000890	1163	R-HEMBB1000890	6672
	HEMBB1000893	F-HEMBB1000893	1164	R-HEMBB1000893	6673
55	HEMBB1000908	F-HEMBB1000908	1165	R-HEMBB1000908	6674

	HEMBB1000910	F-HEMBB1000910	1166	R-HEMBB1000910	6675
	HEMBB1000913	F-HEMBB1000913	1167	R-HEMBB1000913	6676
	HEMBB1000915	F-HEMBB1000915	1168	R-HEMBB1000915	6677
5	HEMBB1000917	F-HEMBB1000917	1169	R-HEMBB1000917	6678
	HEMBB1000927	F-HEMBB1000927	1170	R-HEMBB1000927	6679
	HEMBB1000947	F-HEMBB1000947	1171	R-HEMBB1000947	6680
	HEMBB1000959	F-HEMBB1000959	1172	R-HEMBB1000959	6681
10	HEMBB1000973	F-HEMBB1000973	1173	R-HEMBB1000973	6682
	HEMBB1000975	F-HEMBB1000975	1174	R-HEMBB1000975	6683
	HEMBB1000981	F-HEMBB1000981	1175	R-HEMBB1000981	6684
	HEMBB1000985	F-HEMBB1000985	1176	R-HEMBB1000985	6685
15	HEMBB1000991	F-HEMBB1000991	1177	R-HEMBB1000991	6686
	HEMBB1000996	F-HEMBB1000996	1178	R-HEMBB1000996	6687
	HEMBB1001004	F-HEMBB1001004,	1179	R-HEMBB1001004	6688
	HEMBB1001008	F-HEMBB1001008	1180	R-HEMBB1001008	6689
20	HEMBB1001011	F-HEMBB1001011	1181	R-HEMBB1001011	6690
20	HEMBB1001014	F-HEMBB1001014	1182	R-HEMBB1001014	6691
	HEMBB1001020	F-HEMBB1001020	1183	R-HEMBB1001020	6692
	HEMBB1001024	F-HEMBB1001024	1184	R-HEMBB1001024	6693
	HEMBB1001037	F-HEMBB1001037	1185	R-HEMBB1001037	6694
25	HEMBB1001047	F-HEMBB1001047	1186	R-HEMBB1001047	6695
	HEMBB1001051	F-HEMBB1001051	1187	R-HEMBB1001051	6696
	HEMBB1001056	F-HEMBB1001056	1188	R-HEMBB1001056	6697
	HEMBB1001058	F-HEMBB1001058	1189	R-HEMBB1001058	6698
30	HEMBB1001060	F-HEMBB1001060	1190	R-HEMBB1001060	6699
	HEMBB1001063	F-HEMBB1001063	1191	R-HEMBB1001063	6700
	HEMBB1001068	F-HEMBB1001068	1192	R-HEMBB1001068	6701
	HEMBB1001096	F-HEMBB1001096	1193	R-HEMBB1001096	6702
35	HEMBB1001102	F-HEMBB1001102	1194	R-HEMBB1001102	6703
	HEMBB1001105	F-HEMBB1001105	1195	R-HEMBB1001105	6704
	HEMBB1001112	F-HEMBB1001112	1196		
	HEMBB1001114	F-HEMBB1001114	1197	R-HEMBB1001114	6705
40	HEMBB1001117	F-HEMBB1001117	1198	R-HEMBB1001117	6706
	HEMBB1001119	F-HEMBB1001119	1199	R-HEMBB1001119	6707
	HEMBB1001126	F-HEMBB1001126	1200	R-HEMBB1001126	6708
	HEMBB1001133	F-HEMBB1001133	1201	R-HEMBB1001133	6709
45	HEMBB1001137	F-HEMBB1001137	1202	R-HEMBB1001137	6710
	HEMBB1001142	F-HEMBB1001142	1203	R-HEMBB1001142	6711
	HEMBB1001151	F-HEMBB1001151	1204	R-HEMBB1001151	6712
	HEMBB1001153	F-HEMBB1001153	1205	R-HEMBB1001153	6713
50	HEMBB1001169	F-HEMBB1001169	1206	R-HEMBB1001169	6714
50	HEMBB1001175	F-HEMBB1001175	1207	R-nnnnnnnnnn	6715
	HEMBB1001177	F-HEMBB1001177	1208	R-HEMBB1001177	6716
	HEMBB1001182	F-HEMBB1001182	1209	R-HEMBB1001182	6717
	HEMBB1001199	F-HEMBB1001199	1210	R-HEMBB1001199	6718
55	HEMBB1001208	F-HEMBB1001208	1211	R-HEMBB1001208	6719

	HEMBB1001209	F-HEMBB1001209	1212	R-HEMBB1001209	6720
	HEMBB1001210	F-HEMBB1001210	1213	R-HEMBB1001210	6721
	HEMBB1001218	F-HEMBB1001218	1214	R-HEMBB1001218	6722
5	HEMBB1001221	F-HEMBB1001221	1215	R-HEMBB1001221	6723
	HEMBB1001234	F-HEMBB1001234	1216	R-HEMBB1001234	6724
	HEMBB1001242	F-HEMBB1001242	1217	R-HEMBB1001242	6725
	HEMBB1001249	F-HEMBB1001249	1218	R-HEMBB1001249	6726
10	HEMBB1001253	F-HEMBB1001253	1219	R-HEMBB1001253	6727
	HEMBB1001254	F-HEMBB1001254	1220	R-HEMBB1001254	6728
	HEMBB1001267	F-HEMBB1001267	1221	R-HEMBB1001267	6729
	HEMBB1001271	F-HEMBB1001271	1222	R-HEMBB1001271	6730
15	HEMBB1001282	F-HEMBB1001282	1223	R-HEMBB1001282	6731
70	HEMBB1001288	F-HEMBB1001288	1224	R-HEMBB1001288	6732
	HEMBB1001289	F-HEMBB1001289,	1225	R-HEMBB1001289	6733
	HEMBB1001294	F-HEMBB1001294	1226	R-HEMBB1001294	6734
••	HEMBB1001302	F-HEMBB1001302	1227	R-HEMBB1001302	6735
20	HEMBB1001304	F-HEMBB1001304	1228	R-HEMBB1001304	6736
	HEMBB1001314	F-HEMBB1001314	1229	R-HEMBB1001314	6737
	HEMBB1001315	F-HEMBB1001315	1230	R-HEMBB1001315	6738
	HEMBB1001317	F-HEMBB1001317	1231	R-HEMBB1001317	6739
25	HEMBB1001326	F-HEMBB1001326	1232	R-HEMBB1001326	6740
	HEMBB1001331	F-HEMBB1001331	1233	R-HEMBB1001331	6741
	HEMBB1001335	F-HEMBB1001335	1234	R-HEMBB1001335	6742
	HEMBB1001337	F-HEMBB1001337	1235	R-HEMBB1001337	6743
30	HEMBB1001339	F-HEMBB1001339	1236	R-HEMBB1001339	6744
	HEMBB1001346	F-HEMBB1001346	1237	R-HEMBB1001346	6745
	HEMBB1001348	F-HEMBB1001348	1238	R-HEMBB1001348	6746
	HEMBB1001356	F-HEMBB1001356	1239	R-HEMBB1001356	6747
35	HEMBB1001364	F-HEMBB1001364	1240	R-HEMBB1001364	6748
	HEMBB1001366	F-HEMBB1001366	1241	R-HEMBB1001366	6749
	HEMBB1001367	F-HEMBB1001367	1242	R-HEMBB1001367	6750
	HEMBB1001369	F-HEMBB1001369	1243	R-HEMBB1001369	6751
40	HEMBB1001380	F-HEMBB1001380	1244	R-HEMBB1001380	6752
	HEMBB1001384	F-HEMBB1001384	1245	R-HEMBB1001384	6753
	HEMBB1001387	F-HEMBB1001387	1246	R-HEMBB1001387	6754
	HEMBB1001394	F-HEMBB1001394	1247	R-HEMBB1001394	6755
45	HEMBB1001410	F-HEMBB1001410	1248	R-HEMBB1001410	6756
	HEMBB1001424	F-HEMBB1001424	1249	R-HEMBB1001424	6757
	HEMBB1001426	F-HEMBB1001426	1250	R-HEMBB1001426	6758
	HEMBB1001429	F-HEMBB1001429	1251	R-HEMBB1001429	6759
50	HEMBB1001436	F-HEMBB1001436	1252	R-HEMBB1001436	6760
	HEMBB1001443	F-HEMBB1001443	1253	R-HEMBB1001443	6761
	HEMBB1001449	F-HEMBB1001449	1254	R-HEMBB1001449	6762
	HEMBB1001454	F-HEMBB1001454	1255	R-HEMBB1001454	6763
E E	HEMBB1001458	F-HEMBB1001458	1256	R-HEMBB1001458	6764
55	HEMBB1001463	F-HEMBB1001463	1257	R-HEMBB1001463	6765

	HEMBB1001464	F-HEMBB1001464	1258	R-HEMBB1001464	6766	•
	HEMBB1001482	F-HEMBB1001482	1259	R-HEMBB1001482	6767	
	HEMBB1001500	F-HEMBB1001500	1260	R-HEMBB1001500	6768	
5	HEMBB1001521	F-HEMBB1001521	1261	R-HEMBB1001521	6769	
	HEMBB1001527	F-HEMBB1001527	1262	R-HEMBB1001527	6770	
	HEMBB1001531	F-HEMBB1001531	1263	R-HEMBB1001531	6771	
	HEMBB1001535	F-HEMBB1001535	1264	R-HEMBB1001535	6772	
10	HEMBB1001536	F-HEMBB1001536	1265	R-HEMBB1001536	6773	
	HEMBB1001537	F-HEMBB1001537	1266	R-HEMBB1001537	6774	
	HEMBB1001555	F-HEMBB1001555	1267	R-HEMBB1001555	6775	
	HEMBB1001562	F-HEMBB1001562	1268	R-HEMBB1001562	6776	
15	HEMBB1001564	F-HEMBB1001564	1269	R-HEMBB1001564	6777	
	HEMBB1001565	F-HEMBB1001565	1270	R-HEMBB1001565	6778	
	HEMBB1001585	F-HEMBB1001585,	1271	R-HEMBB1001585	6779	
	HEMBB1001586	F-HEMBB1001586	1272	R-HEMBB1001586	6780	
20	HEMBB1001588	F-HEMBB1001588	1273	R-HEMBB1001588	6781	
20	HEMBB1001603	F-HEMBB1001603	1274	R-HEMBB1001603	6782	
	HEMBB1001618	F-HEMBB1001618	1275	R-HEMBB1001618	6783	
	HEMBB1001619	F-HEMBB1001619	1276	R-HEMBB1001619	6784	
	HEMBB1001630	F-HEMBB1001630	1277	R-HEMBB1001630	6785	
25	HEMBB1001635	F-HEMBB1001635	1278	R-HEMBB1001635	6786	
	HEMBB1001637	F-HEMBB1001637	1279	R-HEMBB1001637	6787	
	HEMBB1001641	F-HEMBB1001641	1280	R-HEMBB1001641	6788	
	HEMBB1001653	F-HEMBB1001653	1281	R-HEMBB1001653	6789	
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	HEMBB1001668	F-HEMBB1001668	1283	R-HEMBB1001668	6791	
	HEMBB1001673	F-HEMBB1001673	1284	R-HEMBB1001673	6792	
	HEMBB1001684	F-HEMBB1001684	1285	R-HEMBB1001684	6793	
35	HEMBB1001685	F-HEMBB1001685	1286	R-HEMBB1001685	6794	
	HEMBB1001695	F-HEMBB1001695	1287	R-HEMBB1001695	6795	
	HEMBB1001704	F-HEMBB1001704	1288	R-HEMBB1001704	6796	
	HEMBB1001706	F-HEMBB1001706	1289	R-HEMBB1001706	6797	
40	HEMBB1001707	F-HEMBB1001707	1290	R-HEMBB1001707	6798	
	HEMBB1001717	F-HEMBB1001717	1291	R-HEMBB1001717	6799	
	HEMBB1001735	F-HEMBB1001735	1292	R-HEMBB1001735	6800	
	HEMBB1001736	F-HEMBB1001736	1293	R-HEMBB1001736	6801	
45	HEMBB1001747	F-HEMBB1001747	1294	R-HEMBB1001747	6802	
	HEMBB1001749	F-HEMBB1001749	1295	R-HEMBB1001749	6803	
	HEMBB1001753	F-HEMBB1001753	1296	R-HEMBB1001753	6804	
	HEMBB1001756	F-HEMBB1001756	1297	R-HEMBB1001756	6805	
50	HEMBB1001760	F-HEMBB1001760	1298	R-HEMBB1001760	6806	
	HEMBB1001762	F-HEMBB1001762	1299	R-HEMBB1001762	6807	
	HEMBB1001785	F-HEMBB1001785	1300	R-HEMBB1001785	6808	
	HEMBB1001797	F-HEMBB1001797	1301	R-HEMBB1001797	6809	
£	HEMBB1001802	F-HEMBB1001802	1302	R-HEMBB1001802	6810	
5 5	HEMBB1001812	F-HEMBB1001812	1303	R-HEMBB1001812	6811	

	HEMBB1001816	F-HEMBB1001816	1304	R-HEMBB1001816	6812
	HEMBB1001831	F-HEMBB1001831	1305	R-HEMBB1001831	6813
	HEMBB1001834	F-HEMBB1001834	1306		
5	HEMBB1001836	F-HEMBB1001836	1307	R-HEMBB1001836	6814
	HEMBB1001839	F-HEMBB1001839	1308	R-HEMBB1001839	6815
	HEMBB1001850	F-HEMBB1001850	1309	R-HEMBB1001850	6816
	HEMBB1001863	F-HEMBB1001863	1310	R-HEMBB1001863	6817
10	HEMBB1001867	F-HEMBB1001867	1311	R-HEMBB1001867	6818
	HEMBB1001868	F-HEMBB1001868	1312	R-HEMBB1001868	6819
	HEMBB1001869	F-HEMBB1001869	1313	R-HEMBB1001869	6820
	HEMBB1001872	F-HEMBB1001872	1314	R-HEMBB1001872	6821
15	HEMBB1001874	F-HEMBB1001874	1315	R-HEMBB1001874	6822
	HEMBB1001875	F-HEMBB1001875	1316	R-HEMBB1001875	6823
	HEMBB1001880	F-HEMBB1001880,	1317	R-HEMBB1001880	6824
	HEMBB1001899	F-HEMBB1001899	1318	R-HEMBB1001899	6825
20	HEMBB1001905	F-HEMBB1001905	1319	R-HEMBB1001905	6826
20	HEMBB1001906	F-HEMBB1001906	1320	R-HEMBB1001906	6827
	HEMBB1001908	F-HEMBB1001908	1321	R-HEMBB1001908	6828
	HEMBB1001910	F-HEMBB1001910	1322	R-HEMBB1001910	6829
0.5	HEMBB1001911	F-HEMBB1001911	1323	R-HEMBB1001911	6830
25	HEMBB1001915	F-HEMBB1001915	1324	R-HEMBB1001915	6831
	HEMBB1001921	F-HEMBB1001921	1325	R-HEMBB1001921	6832
	HEMBB1001922	F-HEMBB1001922	1326	R-HEMBB1001922	6833
	HEMBB1001925	F-HEMBB1001925	1327	R-HEMBB1001925	6834
30	HEMBB1001930	F-HEMBB1001930	1328	R-HEMBB1001930	6835
	HEMBB1001944	F-HEMBB1001944	1329	R-HEMBB1001944	6836
	HEMBB1001945	F-HEMBB1001945	1330	R-HEMBB1001945	6837
	HEMBB1001947	F-HEMBB1001947	1331	R-HEMBB1001947	6838
35	HEMBB1001950	F-HEMBB1001950	1332	R-HEMBB1001950	6839
	HEMBB1001952	F-HEMBB1001952	1333	R-HEMBB1001952	6840
	HEMBB1001953	F-HEMBB1001953	1334	R-HEMBB1001953	6841
	HEMBB1001957	F-HEMBB1001957	1335	R-HEMBB1001957	6842
40	HEMBB1001962	F-HEMBB1001962	1336	R-HEMBB1001962	6843
	HEMBB1001967	F-HEMBB1001967	1337	R-HEMBB1001967	6844
	HEMBB1001973	F-HEMBB1001973	1338	R-HEMBB1001973	6845
	HEMBB1001983	F-HEMBB1001983	1339	R-HEMBB1001983	6846
45	HEMBB1001988	F-HEMBB1001988	1340	R-HEMBB1001988	6847
	HEMBB1001990	F-HEMBB1001990	1341	R-HEMBB1001990	6848
	HEMBB1001996	F-HEMBB1001996	1342	R-HEMBB1001996	6849
	HEMBB1001997	F-HEMBB1001997	1343	R-HEMBB1001997	6850
50	HEMBB1002002	F-HEMBB1002002	1344	R-HEMBB1002002	6851
	HEMBB1002005	F-HEMBB1002005	1345	R-HEMBB1002005	6852
	HEMBB1002009	F-HEMBB1002009	1346	R-HEMBB1002009	6853
	HEMBB1002015	F-HEMBB1002015	1347	R-HEMBB1002015	6854
55	HEMBB1002042	F-HEMBB1002042	1348	R-HEMBB1002042	6855
	HEMBB1002043	F-HEMBB1002043	1349	R-HEMBB1002043	6856

	HEMBB1002044	F-HEMBB1002044	1350	R-HEMBB1002044	6857
	HEMBB1002045	F-HEMBB1002045	1351	R-HEMBB1002045	6858
	HEMBB1002049	F-HEMBB1002049	1352	R-HEMBB1002049	6859
5	HEMBB1002050	F-HEMBB1002050	1353	R-HEMBB1002050	6860
	HEMBB1002068	F-HEMBB1002068	1354	R-HEMBB1002068	6861
	HEMBB1002069	F-HEMBB1002069	1355	R-HEMBB1002069	6862
	HEMBB1002092	F-HEMBB1002092	1356	R-HEMBB1002092	6863
10	HEMBB1002094	F-HEMBB1002094	1357	R-HEMBB1002094	6864
	HEMBB1002115	F-HEMBB1002115	1358	R-HEMBB1002115	6865
	HEMBB1002134	F-HEMBB1002134	1359		
	HEMBB1002139	F-HEMBB1002139	1360	R-HEMBB1002139	6866
15	HEMBB1002142	F-HEMBB1002142	1361	R-HEMBB1002142	6867
15	HEMBB1002152	F-HEMBB1002152	1362	R-HEMBB1002152	6868
	HEMBB1002189	F-HEMBB1002189,	1363	R-HEMBB1002189	6869
	HEMBB1002190	F-HEMBB1002190	1364	R-HEMBB1002190	6870
	HEMBB1002193	F-HEMBB1002193	1365	R-HEMBB1002193	6871
20	HEMBB1002217	F-HEMBB1002217	1366	R-HEMBB1002217	6872
	HEMBB1002218	F-HEMBB1002218	1367	R-HEMBB1002218	6873
	HEMBB1002232	F-HEMBB1002232	1368	R-HEMBB1002232	6874
	HEMBB1002247	F-HEMBB1002247	1369	R-HEMBB1002247	6875
25	HEMBB1002249	F-HEMBB1002249	1370	R-HEMBB1002249	6876
	HEMBB1002254	F-HEMBB1002254	1371	R-HEMBB1002254	6877
	HEMBB1002255	F-HEMBB1002255	1372	R-HEMBB1002255	6878
	HEMBB1002266	F-HEMBB1002266	1373	R-HEMBB1002266	6879
30	HEMBB1002280	F-HEMBB1002280	1374	R-HEMBB1002280	6880
	HEMBB1002300	F-HEMBB1002300	1375	R-HEMBB1002300	6881
	HEMBB1002306	F-HEMBB1002306	1376	R-HEMBB1002306	6882
	HEMBB1002327	F-HEMBB1002327	1377	R-HEMBB1002327	6883
35	HEMBB1002329	F-HEMBB1002329	1378	R-HEMBB1002329	6884
	HEMBB1002340	F-HEMBB1002340	1379	R-HEMBB1002340	6885
	HEMBB1002342	F-HEMBB1002342	1380	R-HEMBB1002342	6886
	HEMBB1002358	F-HEMBB1002358	1381	R-HEMBB1002358	6887
40	HEMBB1002359	F-HEMBB1002359	1382	R-HEMBB1002359	6888
	HEMBB1002364	F-HEMBB1002364	1383	R-HEMBB1002364	6889
	HEMBB1002371	F-HEMBB1002371	1384	R-HEMBB1002371	6890
	HEMBB1002381	F-HEMBB1002381	1385	R-HEMBB1002381	6891
45	HEMBB1002383	F-HEMBB1002383	1386	R-HEMBB1002383	6892
40	HEMBB1002387	F-HEMBB1002387	1387	R-HEMBB1002387	6893
	HEMBB1002409	F-HEMBB1002409	1388		
	HEMBB1002415	F-HEMBB1002415	1389	R-HEMBB1002415	6894
	HEMBB1002425	F-HEMBB1002425	1390	R-HEMBB1002425	6895
50	HEMBB1002442	F-HEMBB1002442	1391	R-HEMBB1002442	6896
	HEMBB1002453	F-HEMBB1002453	1392	R-HEMBB1002453	6897
	HEMBB1002457	F-HEMBB1002457	1393	R-HEMBB1002457	6898
	HEMBB1002458	F-HEMBB1002458	1394	R-HEMBB1002458	6899
55	HEMBB1002477	F-HEMBB1002477	1395	R-HEMBB1002477	6900

	HEMBB1002489	F-HEMBB1002489	1396	R-HEMBB1002489	6901
	HEMBB1002492	F-HEMBB1002492	1397	R-HEMBB1002492	6902
	HEMBB1002495	F-HEMBB1002495	1398	R-HEMBB1002495	6903
5	HEMBB1002502	F-HEMBB1002502	1399	R-HEMBB1002502	6904
	HEMBB1002509	F-HEMBB1002509	1400	R-HEMBB1002509	6905
	HEMBB1002510	F-HEMBB1002510	1401	R-HEMBB1002510	6906
	HEMBB1002520	F-HEMBB1002520	1402	R-HEMBB1002520	6907
10	HEMBB1002522	F-HEMBB1002522	1403	R-HEMBB1002522	6908
	HEMBB1002531	F-HEMBB1002531	1404	R-HEMBB1002531	6909
	HEMBB1002534	F-HEMBB1002534	1405	R-HEMBB1002534	6910
	HEMBB1002545	F-HEMBB1002545	1406	R-HEMBB1002545	6911
15	HEMBB1002550	F-HEMBB1002550	1407	R-HEMBB1002550	6912
	HEMBB1002556	F-HEMBB1002556	1408	R-HEMBB1002556	6913
	HEMBB1002579	F-HEMBB1002579	1409	R-HEMBB1002579	6914
	HEMBB1002582	F-HEMBB1002582	1410	R-HEMBB1002582	6915
20	HEMBB1002590	F-HEMBB1002590	1411	R-HEMBB1002590	6916
20	HEMBB1002596	F-HEMBB1002596	1412	R-HEMBB1002596	6917
	HEMBB1002600	F-HEMBB1002600	1413	R-HEMBB1002600	6918
	HEMBB1002601	F-HEMBB1002601	1414	R-HEMBB1002601	6919
	HEMBB1002603	F-HEMBB1002603	1415	R-HEMBB1002603	6920
25	HEMBB1002607	F-HEMBB1002607	1416	R-HEMBB1002607	6921
	HEMBB1002610	F-HEMBB1002610	1417	R-HEMBB1002610	6922
	HEMBB1002613	F-HEMBB1002613	1418	R-HEMBB1002613	6923
	HEMBB1002614	F-HEMBB1002614	1419	R-HEMBB1002614	6924
30	HEMBB1002617	F-HEMBB1002617	1420	R-HEMBB1002617	6925
	HEMBB1002623	F-HEMBB1002623	1421	R-HEMBB1002623	6926
	HEMBB1002635	F-HEMBB1002635	1422	R-HEMBB1002635	6927
	HEMBB1002664	F-HEMBB1002664	1423	R-HEMBB1002664	6928
35	HEMBB1002677	F-HEMBB1002677	1424	R-HEMBB1002677	6929
	HEMBB1002683	F-HEMBB1002683	1425	R-HEMBB1002683	6930
	HEMBB1002684	F-HEMBB1002684	1426	R-HEMBB1002684	6931
	HEMBB1002686	F-HEMBB1002686	1427	R-HEMBB1002686	6932
40	HEMBB1002692	F-HEMBB1002692	1428	R-HEMBB1002692	6933
	HEMBB1002697	F-HEMBB1002697	1429	R-HEMBB1002697	6934
	HEMBB1002699	F-HEMBB1002699	1430	R-HEMBB1002699	6935
	HEMBB1002702	F-HEMBB1002702	1431	R-HEMBB1002702	6936
45	HEMBB1002705	F-HEMBB1002705	1432	R-HEMBB1002705	6937
	HEMBB1002712	F-HEMBB1002712	1433	R-HEMBB1002712	6938
	MAMMA1000009	F-MAMMA1000009	1434	R-MAMMA1000009	6939
	MAMMA1000019	F-MAMMA1000019	1435	R-MAMMA1000019	6940
50	MAMMA1000020	F-MAMMA1000020	1436	R-MAMMA1000020	6941
50	MAMMA1000025	F-MAMMA1000025	1437	R-MAMMA1000025	6942
	MAMMA1000043	F-MAMMA1000043	1438	R-MAMMA1000043	6943
	MAMMA1000045	F-MAMMA1000045	1439	R-MAMMA1000045	6944
	MAMMA1000055	F-MAMMA1000055	1440	R-MAMMA1000055	6945
55	MAMMA1000057	F-MAMMA1000057	1441	R-MAMMA1000057	6946

	MAMMA1000069	F-MAMMA1000069	1442	R-MAMMA1000069	6947
	MAMMA1000084	F-MAMMA1000084	1443	R-MAMMA1000084	6948
	MAMMA1000085	F-MAMMA1000085	1444	R-MAMMA1000085	6949
5	MAMMA1000092	F-MAMMA1000092	1445	R-MAMMA1000092	6950
	MAMMA1000103	F-MAMMA1000103	1446	R-MAMMA1000103	6951
	MAMMA1000117	F-MAMMA1000117	1447	R-MAMMA1000117	6952
	MAMMA1000129	F-MAMMA1000129	1448	R-MAMMA1000129	6953
10	MAMMA1000133	F-MAMMA1000133	1449	R-MAMMA1000133	6954
	MAMMA1000134	F-MAMMA1000134	1450	R-MAMMA1000134	6955
	MAMMA1000139	F-MAMMA1000139	1451	R-MAMMA1000139	6956
	MAMMA1000143	F-MAMMA1000143	1452	R-MAMMA1000143	6957
15	MAMMA1000155	F-MAMMA1000155	1453	R-MAMMA1000155	6958
10	MAMMA1000163	F-MAMMA1000163	1454	R-MAMMA1000163	6959
	MAMMA1000171	F-MAMMA1000171,	1455	R-MAMMA1000171	6960
	MAMMA1000173	F-MAMMA1000173	1456	R-MAMMA1000173	6961
	MAMMA1000175	F-MAMMA1000175	1457	R-MAMMA1000175	6962
20	MAMMA1000183	F-MAMMA1000183	1458	R-MAMMA1000183	6963
	MAMMA1000198	F-MAMMA1000198	1459	R-MAMMA1000198	6964
	MAMMA1000221	F-MAMMA1000221	1460	R-MAMMA1000221	6965
	MAMMA1000227	F-MAMMA1000227	1461	R-MAMMA1000227	6966
25	MAMMA1000241	F-MAMMA1000241	1462	R-MAMMA1000241	6967
	MAMMA1000251	F-MAMMA1000251	1463	R-MAMMA1000251	6968
	MAMMA1000254	F-MAMMA1000254	1464	R-MAMMA1000254	6969
	MAMMA1000257	F-MAMMA1000257	1465	R-MAMMA1000257	6970
30	MAMMA1000264	F-MAMMA1000264	1466	R-MAMMA1000264	6971
	MAMMA1000266	F-MAMMA1000266	1467	R-MAMMA1000266	6972
	MAMMA1000270	F-MAMMA1000270	1468	R-MAMMA1000270	6973
	MAMMA1000277	F-MAMMA1000277	1469	R-MAMMA1000277	6974
35	MAMMA1000278	F-MAMMA1000278	1470	R-MAMMA1000278	6975
	MAMMA1000279	F-MAMMA1000279	1471	R-MAMMA1000279	6976
	MAMMA1000284	F-MAMMA1000284	1472	R-MAMMA1000284	6977
	MAMMA1000287	F-MAMMA1000287	1473	R-MAMMA1000287	6978
40	MAMMA1000302	F-MAMMA1000302	1474	R-MAMMA1000302	6979
.•	MAMMA1000307	F-MAMMA1000307	1475	R-MAMMA1000307	6980
	MAMMA1000309	F-MAMMA1000309	1476	R-MAMMA1000309	6981
	MAMMA1000312	F-MAMMA1000312	1477	R-MAMMA1000312	6982
45	MAMMA1000313	F-MAMMA1000313	1478	R-MAMMA1000313	6983
45	MAMMA1000331	F-MAMMA1000331	1479	R-MAMMA1000331	6984
	MAMMA1000339	F-MAMMA1000339	1480	R-MAMMA1000339	6985
	MAMMA1000340	F-MAMMA1000340	1481	R-MAMMA1000340	6986
	MAMMA1000348	F-MAMMA1000348	1482	R-MAMMA1000348	6987
50	MAMMA1000356	F-MAMMA1000356	1483	R-MAMMA1000356	6988
	MAMMA1000360	F-MAMMA1000360	1484	R-MAMMA1000360	6989
	MAMMA1000361	F-MAMMA1000361	1485	R-MAMMA1000361	6990
	MAMMA1000372	F-MAMMA1000372	1486	R-MAMMA1000372	6991
55	MAMMA1000385	F-MAMMA1000385	1487	R-MAMMA1000385	6992

	- MAMMA1000388	F-MAMMA1000388	1488	R-MAMMA1000388	6993
	MAMMA1000395	F-MAMMA1000395	1489	R-MAMMA1000395	6994
	MAMMA1000402	F-MAMMA1000402	1490	R-MAMMA1000402	6995
5	MAMMA1000410	F-MAMMA1000410	1491	R-MAMMA1000410	6996
	MAMMA1000413	F-MAMMA1000413	1492	R-MAMMA1000413	6997
	MAMMA1000414	F-MAMMA1000414	1493	R-MAMMA1000414	6998
	MAMMA1000416	F-MAMMA1000416	1494	R-MAMMA1000416	6999
10	MAMMA1000421	F-MAMMA1000421	1495	R-MAMMA1000421	7000
	MAMMA1000422	F-MAMMA1000422	1496	R-MAMMA1000422	7001
	MAMMA1000423	F-MAMMA1000423	1497	R-MAMMA1000423	7002
	MAMMA1000424	F-MAMMA1000424	1498	R-MAMMA1000424	7003
15	MAMMA1000429	F-MAMMA1000429	1499	R-MAMMA1000429	7004
	MAMMA1000431	F-MAMMA1000431	1500	R-MAMMA1000431	7005
	MAMMA1000444	F-MAMMA1000444,	1501	R-MAMMA1000444	7006
	MAMMA1000446	F-MAMMA1000446	1502	R-MAMMA1000446	7007
20	MAMMA1000458	F-MAMMA1000458	1503	R-MAMMA1000458	7008
20	MAMMA1000468	F-MAMMA1000468	1504	R-MAMMA1000468	7009
	MAMMA1000472	F-MAMMA1000472	1505	R-MAMMA1000472	7010
	MAMMA1000478	F-MAMMA1000478	1506	R-MAMMA1000478	7011
	MAMMA1000483	F-MAMMA1000483	1507	R-MAMMA1000483	7012
25	MAMMA1000490	F-MAMMA1000490	1508	R-MAMMA1000490	7013
	MAMMA1000500	F-MAMMA1000500	1509	R-MAMMA1000500	7014
	MAMMA1000501	F-MAMMA1000501	1510	R-MAMMA1000501	7015
	MAMMA1000516	F-MAMMA1000516	1511	R-MAMMA1000516	7016
30	MAMMA1000522	F-MAMMA1000522	1512	R-MAMMA1000522	7017
	MAMMA1000524	F-MAMMA1000524	1513		
	MAMMA1000559	F-MAMMA1000559	1514	R-MAMMA1000559	7018
	MAMMA1000565	F-MAMMA1000565	1515	R-MAMMA1000565	7019
35	MAMMA1000567	F-MAMMA1000567	1516	R-MAMMA1000567	7020
	MAMMA1000576	F-MAMMA1000576	1517	R-MAMMA1000576	7021
	MAMMA1000583	F-MAMMA1000583	1518	R-MAMMA1000583	7022
	MAMMA1000585	F-MAMMA1000585	1519	R-MAMMA1000585	7023
40	MAMMA1000594	F-MAMMA1000594	1520	R-MAMMA1000594	7024
	MAMMA1000597	F-MAMMA1000597	1521	R-MAMMA1000597	7025
	MAMMA1000605	F-MAMMA1000605	1522	R-MAMMA1000605	7026
	MAMMA1000612	F-MAMMA1000612	1523	R-MAMMA1000612	7027
45	MAMMA1000616	F-MAMMA1000616	1524	R-MAMMA1000616	7028
	MAMMA1000621	F-MAMMA1000621	1525	R-MAMMA1000621	7029
	MAMMA1000623	F-MAMMA1000623	1526	R-MAMMA1000623	7030
	MAMMA1000625	F-MAMMA1000625	1527	R-MAMMA1000625	7031
50	MAMMA1000643	F-MAMMA1000643	1528	R-MAMMA1000643	7032
	MAMMA1000664	F-MAMMA1000664	1529	R-MAMMA1000664	7033
	MAMMA1000669	F-MAMMA1000669	1530	R-MAMMA1000669	7034
	MAMMA1000670	F-MAMMA1000670	1531	R-MAMMA1000670	7035
	MAMMA1000672	F-MAMMA1000672	1532	R-MAMMA1000672	7036
55	MAMMA1000684	F-MAMMA1000684	1533	R-MAMMA1000684	7037

	MAMMA1000696	F-MAMMA1000696	1534	R-MAMMA1000696	7038
	MAMMA1000707	F-MAMMA1000707	1535	R-MAMMA1000707	7039
	MAMMA1000713	F-MAMMA1000713	1536	R-MAMMA1000713	7040
5	MAMMA1000714	F-MAMMA1000714	1537	R-MAMMA1000714	7041
	MAMMA1000718	F-MAMMA1000718	1538	R-MAMMA1000718	7042
	MAMMA1000720	F-MAMMA1000720	1539	R-MAMMA1000720	7043
	MAMMA1000723	F-MAMMA1000723	1540	R-MAMMA1000723	7044
10	MAMMA1000731	F-MAMMA1000731	1541	R-MAMMA1000731	7045
	MAMMA1000732	F-MAMMA1000732	1542	R-MAMMA1000732	7046
	MAMMA1000733	F-MAMMA1000733	1543	R-MAMMA1000733	7047
	MAMMA1000734	F-MAMMA1000734	1544	R-MAMMA1000734	7048
45	MAMMA1000734	F-MAMMA1000734	1545	R-MAMMA1000734	7049
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				R-MAMMA1000744	7051
	MAMMA1000746	F-MAMMA1000746	1547		7051
	MAMMA1000752 MAMMA1000760	F-MAMMA1000752	1548	R-MAMMA1000752	7052
20		F-MAMMA1000760	1549	R-MAMMA1000760	
	MAMMA1000761	F-MAMMA1000761	1550	R-MAMMA1000761	7054
	MAMMA1000775	F-MAMMA1000775	1551	R-MAMMA1000775	7055
	MAMMA1000776	F-MAMMA1000776	1552	R-MAMMA1000776	7056
25	MAMMA1000778	F-MAMMA1000778	1553	R-MAMMA1000778	7057
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	MAMMA1000798	F-MAMMA1000798	1555	R-MAMMA1000798	7059
	MAMMA1000802	F-MAMMA1000802	1556	R-MAMMA1000802	7060
30	MAMMA1000824	F-MAMMA1000824	1557	B 1/118/11466664	
30	MAMMA1000831	F-MAMMA1000831	1558	R-MAMMA1000831	7061
	MAMMA1000839	F-MAMMA1000839	1559	R-MAMMA1000839	7062
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35	MAMMA1000843	F-MAMMA1000843	1562	R-MAMMA1000843	7065
	MAMMA1000845	F-MAMMA1000845	1563	R-MAMMA1000845	7066
	MAMMA1000851	F-MAMMA1000851	1564	R-MAMMA1000851	7067
	MAMMA1000855	F-MAMMA1000855	1565	R-MAMMA1000855	7068
40	MAMMA1000856	F-MAMMA1000856	1566	R-MAMMA1000856	7069
	MAMMA1000859	F-MAMMA1000859	1567	•	
	MAMMA1000862	F-MAMMA1000862	1568	R-MAMMA1000862	7070
	MAMMA 1000863	F-MAMMA1000863	1569	R-MAMMA1000863	7071
45	MAMMA1000865	F-MAMMA1000865	1570	R-MAMMA1000865	7072
	MAMMA1000867	F-MAMMA1000867	1571	R-MAMMA1000867	7073
	MAMMA1000875	F-MAMMA1000875	1572	R-MAMMA1000875	7074
	MAMMA1000876	F-MAMMA1000876	1573	R-MAMMA1000876	7075
E0.	MAMMA1000877	F-MAMMA1000877	1574	R-MAMMA1000877	7076
50	MAMMA1000880	F-MAMMA1000880	1575	R-MAMMA1000880	7077
	MAMMA1000883	F-MAMMA1000883	1576	R-MAMMA1000883	7078
	MAMMA1000897	F-MAMMA1000897	1577	R-MAMMA1000897	7079
	MAMMA1000905	F-MAMMA1000905	1578	R-MAMMA1000905	7080
55	MAMMA1000906	F-MAMMA1000906	1579	R-MAMMA1000906	7081

	MAMMA1000908	F-MAMMA1000908	1580	R-MAMMA1000908	7082
	MAMMA1000914	F-MAMMA1000914	1581	R-MAMMA1000914	7083
	MAMMA1000921	F-MAMMA1000921	1582	R-MAMMA1000921	7084
5	MAMMA1000931	F-MAMMA1000931	1583	R-MAMMA1000931	7085
	MAMMA1000940 .	F-MAMMA1000940	1584	R-MAMMA1000940	7086
	MAMMA1000941	F-MAMMA1000941	1585	R-MAMMA1000941	7087
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10	MAMMA1000943	F-MAMMA1000943	1587	R-MAMMA1000943	7089
	MAMMA1000956	F-MAMMA1000956	1588	R-MAMMA1000956	7090
	MAMMA1000957	F-MAMMA1000957	1589	R-MAMMA1000957	7091
	MAMMA1000962	F-MAMMA1000962	1590	R-MAMMA1000962	7092
15	MAMMA1000968	F-MAMMA1000968	1591	R-MAMMA1000968	7093
	MAMMA1000975	F-MAMMA1000975	1592	R-MAMMA1000975	7094
	MAMMA1000979	F-MAMMA1000979,	1593	R-MAMMA1000979	7095
	MAMMA1000987	F-MAMMA1000987	1594	R-MAMMA1000987	7096
20	MAMMA1000998	F-MAMMA1000998	1595	R-MAMMA1000998	7097
20	MAMMA1001003	F-MAMMA1001003	1596	R-MAMMA1001003	7098
	MAMMA1001008	F-MAMMA1001008	1597	R-MAMMA1001008	7099
	MAMMA1001021	F-MAMMA1001021	1598	R-MAMMA1001021	7100
	MAMMA1001024	F-MAMMA1001024	1599	R-MAMMA1001024	7101
25	MAMMA1001030	F-MAMMA1001030	1600	R-MAMMA1001030	7102
	MAMMA1001035	F-MAMMA1001035	1601	R-MAMMA1001035	7103
	MAMMA1001038	F-MAMMA1001038	1602	R-MAMMA1001038	7104
	MAMMA1001041	F-MAMMA1001041	1603	R-nnnnnnnnnnn	7105
30	MAMMA1001050	F-MAMMA1001050	1604	R-MAMMA1001050	7106
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	MAMMA1001067	F-MAMMA1001067	1606	R-MAMMA1001067	7108
	MAMMA1001073	F-MAMMA1001073	1607	R-MAMMA1001073	7109
35	MAMMA1001074	F-MAMMA1001074	1608	R-MAMMA1001074	7110
	MAMMA1001075	F-MAMMA1001075	1609	R-MAMMA1001075	7111
	MAMMA1001078	F-MAMMA1001078	1610	R-MAMMA1001078	7112
	MAMMA1001080	F-MAMMA1001080	1611		
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	MAMMA1001091	F-MAMMA1001091	1613	R-MAMMA1001091	7114
	MAMMA1001092	F-MAMMA1001092	1614	R-MAMMA1001092	7115
	MAMMA1001105	F-MAMMA1001105	1615	R-MAMMA1001105	7116
45	MAMMA1001110	F-MAMMA1001110	1616	R-MAMMA1001110	7117
	MAMMA1001126	F-MAMMA1001126	1617	R-MAMMA1001126	7118
	MAMMA1001133	F-MAMMA1001133	1618	R-MAMMA1001133	7119
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50	MAMMA1001143	F-MAMMA1001143	1620	R-MAMMA1001143	7121
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	MAMMA1001154	F-MAMMA1001154	1622	R-MAMMA1001154	7123
	MAMMA1001161	F-MAMMA1001161	1623	R-MAMMA1001161	7124
55	MAMMA1001162	F-MAMMA1001162	1624	R-MAMMA1001162	7125
00	MAMMA1001181	F-MAMMA1001181	1625	R-MAMMA1001181	7126

	MAMMA1001186	F-MAMMA1001186	1626	R-MAMMA1001186	7127
	MAMMA1001191	F-MAMMA1001191	1627	R-MAMMA1001191	7128
	MAMMA1001198	F-MAMMA1001198	1628	R-MAMMA1001198	7129
5	MAMMA1001202	F-MAMMA1001202	1629	R-MAMMA1001202	7130
	MAMMA1001203	F-MAMMA1001203	1630	R-MAMMA1001203	7131
	MAMMA1001206	F-MAMMA1001206	1631	R-MAMMA1001206	7132
	MAMMA1001215	F-MAMMA1001215	1632	R-MAMMA1001215	7133
10	MAMMA1001220	F-MAMMA1001220	1633	R-MAMMA1001220	7134
	MAMMA1001222	F-MAMMA1001222	1634	R-MAMMA1001222	7135
	MAMMA1001243	F-MAMMA1001243	1635	R-MAMMA1001243	7136
	MAMMA1001244	F-MAMMA1001244	1636	R-MAMMA1001244	7137
15	MAMMA1001249	F-MAMMA1001249	1637	R-MAMMA1001249	7138
13	MAMMA1001256	F-MAMMA1001256	1638	R-MAMMA1001256	7139
	MAMMA1001259	F-MAMMA1001259	1639	R-MAMMA1001259	7140
	MAMMA1001260	F-MAMMA1001260	1640	R-MAMMA1001260	7141
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20	MAMMA1001271	F-MAMMA1001271	1642	R-MAMMA1001271	7143
	MAMMA1001274	F-MAMMA1001274	1643	R-MAMMA1001274	7144
	MAMMA1001280	F-MAMMA1001280	1644	R-MAMMA1001280	7145
	MAMMA1001292	F-MAMMA1001292	1645	R-MAMMA1001292	7146
25	MAMMA1001296	F-MAMMA1001296	1646	R-MAMMA1001296	7147
	MAMMA1001298	F-MAMMA1001298	1647	R-MAMMA1001298	7148
	MAMMA1001305	F-MAMMA1001305	1648	R-MAMMA1001305	7149
	MAMMA1001322	F-MAMMA1001322	1649	R-MAMMA1001322	7150
30	MAMMA1001324	F-MAMMA1001324	1650	R-MAMMA1001324	7151
	MAMMA1001330	F-MAMMA1001330	1651	R-MAMMA1001330	7152
	MAMMA1001341	F-MAMMA1001341	1652	R-MAMMA1001341	7153
	MAMMA1001343	F-MAMMA1001343	1653	R-MAMMA1001343	7154
35	MAMMA1001346	F-MAMMA1001346	1654	R-MAMMA1001346	7155
	MAMMA1001383	F-MAMMA1001383	1655	R-MAMMA1001383	7156
	MAMMA1001388	F-MAMMA1001388	1656	R-MAMMA1001388	7157
	MAMMA1001397	F-MAMMA1001397	1657	R-MAMMA1001397	7158
40	MAMMA1001408	F-MAMMA1001408	1658	R-MAMMA1001408	7159
40	MAMMA1001411	F-MAMMA1001411	1659	R-MAMMA1001411	7160
	MAMMA1001419	F-MAMMA1001419	1660	R-MAMMA1001419	7161
	MAMMA1001420	F-MAMMA1001420	1661	R-MAMMA1001420	7162
	MAMMA1001435	F-MAMMA1001435	1662	R-MAMMA1001435	7163
45	MAMMA1001442	F-MAMMA1001442	1663	R-MAMMA1001442	7164
	MAMMA1001446	F-MAMMA1001446	1664	R-MAMMA1001446	7165
	MAMMA1001452	F-MAMMA1001452	1665	R-MAMMA1001452	7166
	MAMMA1001465	F-MAMMA1001465	1666	R-MAMMA1001465	7167
50	MAMMA1001476	F-MAMMA1001476	1667	R-MAMMA1001476	7168
	MAMMA1001487	F-MAMMA1001487	1668	R-MAMMA1001487	7169
	MAMMA1001501	F-MAMMA1001501	1669	R-MAMMA1001501	7170
	MAMMA1001502	F-MAMMA1001502	1670	R-MAMMA1001502	7171
55	MAMMA1001510	F-MAMMA1001510	1671	R-MAMMA1001510	7172

	MAMMA1001522	F-MAMMA1001522	1672	R-MAMMA1001522	7173
	MAMMA1001547	F-MAMMA1001547	1673	R-MAMMA1001547	7174
	MAMMA1001551	F-MAMMA1001551	1674	R-MAMMA1001551	7175
5	MAMMA1001575	F-MAMMA1001575	1675	R-MAMMA1001575	7176
	MAMMA1001576	F-MAMMA1001576	1676	R-MAMMA1001576	7177
	MAMMA1001590	F-MAMMA1001590	1677	R-MAMMA1001590	7178
	MAMMA1001600	F-MAMMA1001600	1678	R-MAMMA1001600	7179
10	MAMMA1001604	F-MAMMA1001604	1679	R-MAMMA1001604	7180
	MAMMA1001606	F-MAMMA1001606	1680	R-MAMMA1001606	7181
	MAMMA1001620	F-MAMMA1001620	1681	R-MAMMA1001620	7182
	MAMMA1001627	F-MAMMA1001627	1682	R-MAMMA1001627	7183
45	MAMMA1001630	F-MAMMA1001630	1683	R-MAMMA1001630	7184
15	MAMMA1001633	F-MAMMA1001633	1684	R-MAMMA1001633	7185
	MAMMA1001635	F-MAMMA1001635,	1685	R-MAMMA1001635	7186
	MAMMA1001649	F-MAMMA1001649	1686	R-MAMMA1001649	7187
	MAMMA1001654	F-MAMMA1001654	1687		
0	MAMMA1001663	F-MAMMA1001663	1688	R-MAMMA1001663	7188
	MAMMA1001670	F-MAMMA1001670	1689	R-MAMMA1001670	7189
	MAMMA1001671	F-MAMMA1001671	1690	R-MAMMA1001671	7190
	MAMMA1001679	F-MAMMA1001679	1691	R-MAMMA1001679	7191
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	MAMMA1001686	F-MAMMA1001686	1693	R-MAMMA1001686	7193
	MAMMA1001692	F-MAMMA1001692	1694	R-MAMMA1001692	7194
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30	MAMMA1001715	F-MAMMA1001715	1696	R-MAMMA1001715	7196
	MAMMA1001730	F-MAMMA1001730	1697	R-MAMMA1001730	7197
	MAMMA1001735	F-MAMMA1001735	1698	R-MAMMA1001735	7198
	MAMMA1001740	F-MAMMA1001740	1699	R-MAMMA1001740	7199
35	MAMMA1001743	F-MAMMA1001743	1700	R-MAMMA1001743	7200
	MAMMA1001744	F-MAMMA1001744	1701	R-MAMMA1001744	7201
	MAMMA1001745	F-MAMMA1001745	1702	R-MAMMA1001745	7202
	MAMMA1001751	F-MAMMA1001751	1703	R-MAMMA1001751	7203
40	MAMMA1001754	F-MAMMA1001754	1704	R-MAMMA1001754	7204
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	MAMMA1001783	F-MAMMA1001783	1711	R-MAMMA1001783	7211
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	MAMMA1001806	F-MAMMA1001806	1715	R-MAMMA1001806	7215
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	MAMMA1001820	F-MAMMA1001820	1720	R-MAMMA1001820	7220
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	MAMMA1001874	F-MAMMA1001874	1730	R-MAMMA1001874	7230
	MAMMA1001878	F-MAMMA1001878	1731	R-MAMMA1001878	7231
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	MAMMA1002032	F-MAMMA1002032	1744	R-MAMMA1002032	7244
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35	MAMMA1002041	F-MAMMA1002041	1746	R-MAMMA1002041	7246
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	MAMMA1002047	F-MAMMA1002047	1748	R-MAMMA1002047	7248
	MAMMA1002056	F-MAMMA1002056	1749	R-MAMMA1002056	7249
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	MAMMA1002082	F-MAMMA1002082	1753	R-MAMMA1002082	7253
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	MAMMA1002108	F-MAMMA1002108	1756	R-MAMMA1002108	7256
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50	MAMMA1002132	F-MAMMA1002132	1759	R-MAMMA1002132	7259
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	MAMMA1002145	F-MAMMA1002145	1762	R-MAMMA1002145	7262
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	MAMMA1002156	F-MAMMA1002156	1765	R-MAMMA1002156	7265
	MAMMA1002158	F-MAMMA1002158	1766	R-MAMMA1002158	7266
5	MAMMA1002170	F-MAMMA1002170	1767	R-MAMMA1002170	7267
	MAMMA1002174	F-MAMMA1002174	1768	R-MAMMA1002174	7268
	MAMMA1002198	F-MAMMA1002198	1769	R-MAMMA1002198	7269
	MAMMA1002209	F-MAMMA1002209	1770	R-MAMMA1002209	7270
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	MAMMA1002230	F-MAMMA1002230	1773	R-MAMMA1002230	7273
	MAMMA1002236	F-MAMMA1002236	1774	R-MAMMA1002236	7274
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	MAMMA1002268	F-MAMMA1002268	1778	R-MAMMA1002268	7278
	MAMMA1002269	F-MAMMA1002269	1779	R-MAMMA1002269	7279
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	MAMMA1002292	F-MAMMA1002292	1781	R-MAMMA1002292	7281
	MAMMA1002293	F-MAMMA1002293	1782	R-MAMMA1002293	7282
	MAMMA1002294	F-MAMMA1002294	1783	R-MAMMA1002294	7283
25	MAMMA1002297	F-MAMMA1002297	1784	R-MAMMA1002297	7284
	MAMMA1002298	F-MAMMA1002298	1785	R-MAMMA1002298	7285
	MAMMA1002299	F-MAMMA1002299	1786	R-MAMMA1002299	7286
	MAMMA1002308	F-MAMMA1002308	1787	R-MAMMA1002308	7287
30	MAMMA1002310	F-MAMMA1002310	1788	R-MAMMA1002310	7288
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35	MAMMA1002319	F-MAMMA1002319	1792	R-MAMMA1002319	7292
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	MAMMA1002329	F-MAMMA1002329	1794	R-MAMMA1002329	7294
	MAMMA1002332	F-MAMMA1002332	1795	R-MAMMA1002332	7295
40	MAMMA1002333	F-MAMMA1002333	1796	R-MAMMA1002333	7296
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	MAMMA1002347	F-MAMMA1002347	1798	R-MAMMA1002347	7298
	MAMMA1002351	F-MAMMA1002351	1799	R-MAMMA1002351	7299
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70	MAMMA1002353	F-MAMMA1002353	1801	R-MAMMA1002353	7301
	MAMMA1002355	F-MAMMA1002355	1802	R-MAMMA1002355	7302
	MAMMA1002356	F-MAMMA1002356	1803	R-MAMMA1002356	7303
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	MAMMA1002361	F-MAMMA1002361	1806	R-MAMMA1002361	7306
	MAMMA1002362 MAMMA1002380	F-MAMMA1002362 F-MAMMA1002380	1807 1808	R-MAMMA1002362 R-MAMMA1002380	7307
	MAMMA1002384	F-MAMMA1002384	1809	R-MAMMA1002384	7308 7309
55	manua 1002364	1. Mramma1007204	1003	A MARIANTAL TOUGOOG	1309

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15	MAMMA1002475	F-MAMMA1002475	1822	R-MAMMA1002475	7322
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	MAMMA1002485	F-MAMMA1002485	1824	R-MAMMA1002485	7324
	MAMMA1002494	F-MAMMA1002494	1825	R-MAMMA1002494	7325
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	MAMMA1002524	F-MAMMA1002524	1827	R-MAMMA1002524	7327
	MAMMA1002530	F-MAMMA1002530	1828	R-MAMMA1002530	7328
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25	MAMMA1002554	F-MAMMA1002554	1830	R-MAMMA1002554	7330
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	MAMMA1002566	F-MAMMA1002566	1832	R-MAMMA1002566	7332
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30	MAMMA1002573	F-MAMMA1002573	1834	R-MAMMA1002573	7334
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	MAMMA1002597	F-MAMMA1002597	1837	R-MAMMA1002597	7337
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35	MAMMA1002603	F-MAMMA1002603	1839	R-MAMMA1002603	7339
	MAMMA1002612	F-MAMMA1002612	1840	R-MAMMA1002612	7340
	MAMMA1002617	F-MAMMA1002617	1841	R-MAMMA1002617	7341
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40	MAMMA1002619	F-MAMMA1002619	1843	R-MAMMA1002619	7343
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	MAMMA1002637	F-MAMMA1002637	1849	R-MAMMA1002637	7349
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	MAMMA1002662		1852	R-MAMMA1002662	7352
		F-MAMMA1002662	1853	R-MAMMA1002665	7354
	MAMMA1002665	F-MAMMA1002665	1854	R-MAMMA1002651	7354
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70	MAMMA1002721	F-MAMMA1002721	1864	R-MAMMA1002721	7364
	MAMMA1002727	F-MAMMA1002727	1865	R-MAMMA1002727	7365
	MAMMA1002728	F-MAMMA1002728	1866	R-MAMMA1002728	7366
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15	MAMMA1002746	F-MAMMA1002746	1868	R-MAMMA1002746	7368
	MAMMA1002748	F-MAMMA1002748,	1869	R-MAMMA1002748	7369
	MAMMA1002754	F-MAMMA1002754	1870	R-MAMMA1002754	7370
	MAMMA1002754	F-MAMMA1002758	1871	R-MAMMA1002754	7371
20	MAMMA1002764	F-MAMMA1002764	1872	R-MAMMA1002764	7372
	MAMMA1002765	F-MAMMA1002765	1873	R-MAMMA1002765	7373
	MAMMA1002769	F-MAMMA1002769	1874	R-MAMMA1002769	7374
	MAMMA1002775	F-MAMMA1002775	1875	N MARKETTOOL 100	
25	MAMMA1002780	F-MAMMA1002780	1876	R-MAMMA1002780	7375
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	MAMMA1002844	F-MAMMA1002844	1887	R-MAMMA1002844	7386
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40	MAMMA1002868	F-MAMMA1002868	1889	R-MAMMA1002868	7388
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50	MAMMA1002892	F-MAMMA1002892	1897	R-MAMMA1002892	7395
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10	MAMMA1002973	F-MAMMA1002973	1909	R-MAMMA1002973	7406
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	MAMMA1002987	F-MAMMA1002987	1911	R-MAMMA1002987	7408
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15	MAMMA1003004	F-MAMMA1003004	1913	R-MAMMA1003004	7410
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	MAMMA1003031	F-MAMMA1003031	1920	R-MAMMA1003031	7416
25	MAMMA1003035	F-MAMMA1003035	1921	R-MAMMA1003035	7417
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	MAMMA1003040	F-MAMMA1003040	1923	R-MAMMA1003040	7419
	MAMMA1003044	F-MAMMA1003044	1924	R-MAMMA1003044	7420
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	MAMMA1003055	F-MAMMA1003055	1927	R-MAMMA1003055	7423
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	MAMMA1003057	F-MAMMA1003057	1929	R-MAMMA1003057	7425
35	MAMMA1003066	F-MAMMA1003066	1930	R-MAMMA1003066	7426
	MAMMA1003089	F-MAMMA1003089	1931	R-MAMMA1003089	7427
	MAMMA1003099	F-MAMMA1003099	1932	R-MAMMA1003099	7428
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40	MAMMA1003113	F-MAMMA1003113	1934	R-MAMMA1003113	7430
	MAMMA1003127	F-MAMMA1003127	1935	R-MAMMA1003127	7431
	MAMMA1003135	F-MAMMA1003135	1936	R-MAMMA1003135	7432
	MAMMA1003140	F-MAMMA1003140	1937	R-MAMMA1003140	7433
45	MAMMA1003146	F-MAMMA1003146	1938	R-MAMMA1003146	7434
	MAMMA1003150	F-MAMMA1003150	1939	R-nnnnnnnnnnn	7435
	MAMMA1003166	F-MAMMA1003166	1940	R-MAMMA1003166	7436
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	NT2RM1000032	F-NT2RM1000032	1943		
	NT2RM1000035	F-NT2RM1000035	1944		
	NT2RM1000037	F-NT2RM1000037	1945		
55	NT2RM1000039	F-NT2RM1000039	1946		
	NT2RM1000055	F-NT2RM1000055	1947		

	NT2RM1000059	F-NT2RM1000059	1948
	NT2RM1000062	F-NT2RM1000062	1949
	NT2RM1000080	F-NT2RM1000080	1950
5	NT2RM1000086	F-NT2RM1000086	1951
	NT2RM1000092	F-NT2RM1000092	1952
	NT2RM1000118	F-NT2RM1000118	1953
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10	NT2RM1000127	F-NT2RM1000127	1955
	NT2RM1000131	F-NT2RM1000131	1956
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35	NT2RM2000407	F-NT2RM2000407	2068
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45	NT2RM2000522	F-NT2RM2000522	2076
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15	NT2RM2000714	F-NT2RM2000714	2097
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50	NT2RM2001319	F-NT2RM2001319	2127
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55	NT2RM2001370	F-NT2RM2001370	2131

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15	NT2RM2001592	F-NT2RM2001592	2143
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	NT2RM2001641	F-NT2RM2001641	2149
	NT2RM2001648	F-NT2RM2001648	2150
	NT2RM2001652	F-NT2RM2001652	2151
25	NT2RM2001659	F-NT2RM2001659	2152
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35	NT2RM2001695	F-NT2RM2001695	2160
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45	NT2RM2001727	F-NT2RM2001727	2169
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	NT2RM2001771	F-NT2RM2001771	2175
	NT2RM2001782	F-NT2RM2001782	2176
	NT2RM2001784	F-NT2RM2001784	2177
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	NT2RM2001855	F-NT2RM2001855	2187		
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	NT2RM2001903	F-NT2RM2001903	2192		
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20	NT2RM2001935	F-NT2RM2001935	2194		
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30	NT2RM2002014	F-NT2RM2002014	2203		
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	NT2RM2002049	F-NT2RM2002049	2205	•	
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	NT2RM4000027	F-NT2RM4000027	2217	R-NT2RM4000027	7439
	NT2RM4000030	F-NT2RM4000030	2218	R-NT2RM4000030	7440
50	NT2RM4000046	F-NT2RM4000046	2219	R-NT2RM4000046	7441
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	NT2RM4000086	F-NT2RM4000086	2222	R-NT2RM4000086	7444
	NT2RM4000104	F-NT2RM4000104	2223	R-NT2RM4000104	7445
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	NT2RM4000155	F-NT2RM4000155	2225	R-NT2RM4000155	7447
	NT2RM4000156	F-NT2RM4000156	2226	R-NT2RM4000156	7448
5	NT2RM4000167	F-NT2RM4000167	2227	R-nnnnnnnnnnn	7449
	NT2RM4000169	F-NT2RM4000169	2228	R-NT2RM4000169	7450
	NT2RM4000191	F-NT2RM4000191	2229	R-NT2RM4000191	7451
	NT2RM4000197	F-NT2RM4000197	2230	R-NT2RM4000197	7452
10	NT2RM4000199	F-NT2RM4000199	2231	R-NT2RM4000199	7453
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	NT2RM4000202	F-NT2RM4000202	2233	R-NT2RM4000202	7455
	NT2RM4000210	F-NT2RM4000210	2234	R-NT2RM4000210	7456
	NT2RM4000215	F-NT2RM4000215	2235	R-NT2RM4000215	7457
15	NT2RM4000229	F-NT2RM4000229	2236	R-nnnnnnnnnn	7458
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	NT2RM4000290	F-NT2RM4000290	2241	R-NT2RM4000290	7463
	NT2RM4000324	F-NT2RM4000324	2242	R-NT2RM4000324	7464
	NT2RM4000327	F-NT2RM4000327	2243	R-NT2RM4000327	7465
25	NT2RM4000344	F-NT2RM4000344	2244	R-NT2RM4000344	7466
	NT2RM4000349	F-NT2RM4000349	2245	R-NT2RM4000349	7467
	NT2RM4000354	F-NT2RM4000354	2246	R-NT2RM4000354	7468
	NT2RM4000356	F-NT2RM4000356	2247	R-NT2RM4000356	7469
30	NT2RM4000366	F-NT2RM4000366	2248	R-NT2RM4000366	7470
	NT2RM4000368	F-NT2RM4000368	2249	R-NT2RM4000368	7471
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	NT2RM4000395	F-NT2RM4000395	2251	R-NT2RM4000395	7473
	NT2RM4000414	F-NT2RM4000414	2252	R-NT2RM4000414	7474
35	NT2RM4000421	F-NT2RM4000421	2253	R-NT2RM4000421	7475
	NT2RM4000425	F-NT2RM4000425	2254	R-NT2RM4000425	7476
	NT2RM4000433	F-NT2RM4000433	2255	R-NT2RM4000433	7477
	NT2RM4000457	F-NT2RM4000457	2256	R-NT2RM4000457	7478
40	NT2RM4000471	F-NT2RM4000471	2257	R-NT2RM4000471	7479
	NT2RM4000486	F-NT2RM4000486	2258	R-NT2RM4000486	7480
	NT2RM4000496	F-NT2RM4000496	2259	R-NT2RM4000496	7481
	NT2RM4000511	F-NT2RM4000511	2260	R-NT2RM4000511	7482
45	NT2RM4000514	F-NT2RM4000514	2261	R-NT2RM4000514	7483
	NT2RM4000515	F-NT2RM4000515	2262	R-nnnnnnnnnnn	7484
	NT2RM4000520	F-NT2RM4000520	2263	R-NT2RM4000520	7485
	NT2RM4000531	F-NT2RM4000531	2264	R-NT2RM4000531	7486
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	NT2RM4000585	F-NT2RM4000585	2267	R-NT2RM4000585	7489
	NT2RM4000590	F-NT2RM4000590	2268	R-NT2RM4000590	7490
	NT2RM4000595	F-NT2RM4000595	2269	R-NT2RM4000595	7491
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	NT2RM4000616	F-NT2RM4000616	2272	R-NT2RM4000616	7494
5	NT2RM4000674	F-NT2RM4000674	2273	R-NT2RM4000674	7495
	NT2RM4000689	F-NT2RM4000689	2274	R-NT2RM4000689	7496
	NT2RM4000698	F-NT2RM4000698	2275	R-NT2RM4000698	7497
	NT2RM4000700	F-NT2RM4000700	2276	R-nnnnnnnnnnn	7498
10	NT2RM4000712	F-NT2RM4000712	2277	R-NT2RM4000712	7499
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	NT2RM4000733	F-NT2RM4000733	2279	R-NT2RM4000733	7501
	NT2RM4000734	F-NT2RM4000734	2280	R-NT2RM4000734	7502
	NT2RM4000741	F-NT2RM4000741	2281	R-NT2RM4000741	7503
15	NT2RM4000751	F-NT2RM4000751	2282	R-NT2RM4000751	7504
	NT2RM4000764	F-NT2RM4000764,	2283	R-NT2RM4000764	7505
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	NT2RM4000779	F-NT2RM4000779	2285	R-NT2RM4000779	7507
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25	NT2RM4000798	F-NT2RM4000798	2290	R-NT2RM4000798	7512
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	NT2RM4000820	F-NT2RM4000820	2292	R-NT2RM4000820	7514
	NT2RM4000833	F-NT2RM4000833	2293	R-NT2RM4000833	7515
30	NT2RM4000848	F-NT2RM4000848	2294	R-NT2RM4000848	7516
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	NT2RM4000855	F-NT2RM4000855	2296	R-NT2RM4000855	7518
	NT2RM4000887	F-NT2RM4000887	2297	R-nnnnnnnnnnn	7519
	NT2RM4000895	F-NT2RM4000895	2298	R-NT2RM4000895	7520
35	NT2RM4000950	F-NT2RM4000950	2299	R-NT2RM4000950	7521
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	NT2RM4000979	F-NT2RM4000979	2301	R-NT2RM4000979	7523
	NT2RM4000996	F-NT2RM4000996	2302	R-NT2RM4000996	7524
40	NT2RM4001002	F-NT2RM4001002	2303	R-NT2RM4001002	7525
	NT2RM4001016	F-NT2RM4001016	2304	R-NT2RM4001016	7526
	NT2RM4001032	F-NT2RM4001032	2305	R-NT2RM4001032	7527
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	NT2RM4001092	F-NT2RM4001092	2309	R-NT2RM4001092	7531
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	NT2RM4001155	F-NT2RM4001155	2313	R-NT2RM4001155	7535
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	NT2RM4001217	F-NT2RM4001217	2320	R-NT2RM4001217	7542
	NT2RM4001256	F-NT2RM4001256	2321	R-NT2RM4001256	7543
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	NT2RM4001371	F-NT2RM4001371	2330	R-NT2RM4001371	7552
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	NT2RM4001650	F-NT2RM4001650	2356	R-NT2RM4001650	7578
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	NT2RM4001841	F-NT2RM4001841	2376	R-NT2RM4001841	7597
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	NT2RM4001940	F-NT2RM4001940	2387	R-NT2RM4001940	7608
	NT2RM4001953	F-NT2RM4001953	2388	R-NT2RM4001953	7609 7610
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	NT2RM4002055		2399		
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10	NT2RM4002194	F-NT2RM4002194	2415	R-NT2RM4002194	7635
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	NT2RM4002301	F-NT2RM4002301	2426	R-NT2RM4002301	7646
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25	NT2RM4002339	F-NT2RM4002339	2428	R-nnnnnnnnnnn	7648
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	NT2RM4002390	F-NT2RM4002390	2433	R-NT2RM4002390	7653
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	NT2RM4002452	F-NT2RM4002452	2438	R-NT2RM4002452	7657
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	NT2RM4002527	F-NT2RM4002527	2446	R-nananananan	7665
	NT2RM4002532	F-NT2RM4002532	2447	R-NT2RM4002532	7666
	NT2RM4002534	F-NT2RM4002534	2448	R-NT2RM4002534	7667
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50	NT2RM4002565	F-NT2RM4002565	2450		
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	NT2RM4002571	F-NT2RM4002571	2452	R-NT2RM4002571	7669
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	NT2RP1000063	F-NT2RP1000063	2459		
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	NT2RP1000112	F-NT2RP1000112	2463		
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	NT2RP1000513	F-NT2RP1000513	2493		
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25	NT2RP1000856	F-NT2RP1000856	2520
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35	NT2RP2000007	F-NT2RP2000007	2574		
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	NT2RP2000097	F-NT2RP2000097	2589	R-NT2RP2000097	7687
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	NT2RP2000153	F-NT2RP2000153	2597	R-NT2RP2000153	7695
	NT2RP2000157	F-NT2RP2000157	2598	R-NT2RP2000157	7696
10	NT2RP2000161	F-NT2RP2000161	2599	R-NT2RP2000161	7697
	NT2RP2000173	F-NT2RP2000173	2600		
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	NT2RP2000183	F-NT2RP2000183	2602	R-NT2RP2000183	7699
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	NT2RP2000208	F-NT2RP2000208	2605		
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20	NT2RP2000232	F-NT2RP2000232	2607	R-NT2RP2000232	7703
20	NT2RP2000233	F-NT2RP2000233	2608	R-NT2RP2000233	7704
	NT2RP2000239	F-NT2RP2000239	2609	R-NT2RP2000239	7705
	NT2RP2000248	F-NT2RP2000248	2610	R-NT2RP2000248	7706
	NT2RP2000257	F-NT2RP2000257	2611	R-NT2RP2000257	7707
25	NT2RP2000258	F-NT2RP2000258	2612	R-NT2RP2000258	7708
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	NT2RP2000274	F-NT2RP2000274	2614	R-NT2RP2000274	7710
	NT2RP2000283	F-NT2RP2000283	2615		
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	NT2RP2000289	F-NT2RP2000289	2617	R-NT2RP2000289	7712
	NT2RP2000297	F-NT2RP2000297	2618	R-NT2RP2000297	7713
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	NT2RP2000327	F-NT2RP2000327	2621	R-NT2RP2000327	7716
	NT2RP2000328	F-NT2RP2000328	2622		
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40	NT2RP2000337	F-NT2RP2000337	2624	R-NT2RP2000337	7718
	NT2RP2000346	F-NT2RP2000346	2625	R-NT2RP2000346	7719
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	NT2RP2000412	F-NT2RP2000412	2627		
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••	NT2RP2000459	F-NT2RP2000459	2633	R-NT2RP2000459	7726
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	NT2RP2000617	F-NT2RP2000617	2640	R-NT2RP2000617	7733
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	NT2RP2000658	F-NT2RP2000658	2644	R-NT2RP2000658	7737
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	NT2RP2000704	F-NT2RP2000704	2647		
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	NT2RP2000764	F-NT2RP2000764	2652	R-NT2RP2000764	7744
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	NT2RP2000814	F-NT2RP2000814	2655	R-nnnnnnnnnnn	7747
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	NT2RP2000932	F-NT2RP2000932	2665		
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	NT2RP2001044	F-NT2RP2001044	2673	R-NT2RP2001044	7764
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	NT2RP2001070	F-NT2RP2001070	2676	R-NT2RP2001070	7766
	NT2RP2001081	F-NT2RP2001081	2677		
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	NT2RP2001174	F-NT2RP2001174	2685	R-NT2RP2001174	7774
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	NT2RP2001277	F-NT2RP2001277	2692	R-NT2RP2001277	7781
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	NT2RP2001378	F-NT2RP2001378	2700	R-NT2RP2001378	7788
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	NT2RP2001423	F-NT2RP2001423	2706	R-NT2RP2001423	7794
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	NT2RP2001536	F-NT2RP2001536	2718	R-NT2RP2001536	7806
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	NT2RP2001597	F-NT2RP2001597	2723	R-NT2RP2001597	7811
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	NT2RP2001634	F-NT2RP2001634	2727		
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5	NT2RP2001699	F-NT2RP2001699	2733	R-NT2RP2001699	7818
	NT2RP2001720	F-NT2RP2001720	2734	R-NT2RP2001720	7819
	NT2RP2001721	F-NT2RP2001721	2735	R-NT2RP2001721	7820
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	NT2RP2001762	F-NT2RP2001762	2738	R-NT2RP2001762	7823
	NT2RP2001813	F-NT2RP2001813	2739	R-NT2RP2001813	7824
	NT2RP2001839	F-NT2RP2001839	2740		
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.•	NT2RP2001869	F-NT2RP2001869	2742	R-NT2RP2001869	7826
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	NT2RP2001907	F-NT2RP2001907	2747	R-NT2RP2001907	7830
	NT2RP2001926	F-NT2RP2001926	2748	R-NT2RP2001926	7831
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25	NT2RP2001943	F-NT2RP2001943	2750	R-NT2RP2001943	7833
	NT2RP2001946	F-NT2RP2001946	2751	R-NT2RP2001946	7834
	NT2RP2001947	F-NT2RP2001947	2752	R-NT2RP2001947	7835
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	NT2RP2001985	F-NT2RP2001985	2755	R-NT2RP2001985	7838
	NT2RP2001991	F-NT2RP2001991	2756		
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35	NT2RP2002032	F-NT2RP2002032	2758	R-NT2RP2002032	7840
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	NT2RP2002066	F-NT2RP2002066	2764	R-NT2RP2002066	7846
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		NT2RP2002193	F-NT2RP2002193	2777	R-NT2RP2002193	7858
		NT2RP2002208	F-NT2RP2002208	2778	R-NT2RP2002208	7859
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		NT2RP2002235	F-NT2RP2002235	2781		
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		NT2RP2002270	F-NT2RP2002270	2785	R-NT2RP2002270	7865
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	5	NT2RP2002312	F-NT2RP2002312	2787	R-NT2RP2002312	7867
,	3	NT2RP2002316	F-NT2RP2002316	2788	R-NT2RP2002316	7868
		NT2RP2002325	F-NT2RP2002325,	2789	R-NT2RP2002325	7869
		NT2RP2002333	F-NT2RP2002333	2790	R-NT2RP2002333	7870
		NT2RP2002373	F-NT2RP2002373	2791		
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		NT2RP2002394	F-NT2RP2002394	2793	R-NT2RP2002394	7872
		NT2RP2002408	F-NT2RP2002408	2794	R-NT2RP2002408	7873
		NT2RP2002426	F-NT2RP2002426	2795	R-NT2RP2002426	7874
2	25	NT2RP2002439	F-NT2RP2002439	2796	R-NT2RP2002439	7875
		NT2RP2002442	F-NT2RP2002442	2797	•	
		NT2RP2002457	F-NT2RP2002457	2798	R-NT2RP2002457	7876
		NT2RP2002464	F-NT2RP2002464	2799	R-NT2RP2002464	7877
3	10	NT2RP2002475	F-NT2RP2002475	2800	R-NT2RP2002475	7878
		NT2RP2002479	F-NT2RP2002479	2801	R-nnnnnnnnnnn	7879
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3	15	NT2RP2002504	F-NT2RP2002504	2804	R-NT2RP2002504	7882
		NT2RP2002520	F-NT2RP2002520	2805	R-NT2RP2002520	7883
		NT2RP2002537	F-NT2RP2002537	2806	R-NT2RP2002537	7884
		NT2RP2002546	F-NT2RP2002546	2807	R-NT2RP2002546	7885
	10	NT2RP2002549	F-NT2RP2002549	2808	R-NT2RP2002549	7886
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		NT2RP2002595	F-NT2RP2002595	2810	R-NT2RP2002595	7888
		NT2RP2002606	F-NT2RP2002606	2811	R-NT2RP2002606	7889
		NT2RP2002609	F-NT2RP2002609	2812	R-NT2RP2002609	7890
4	15	NT2RP2002618	F-NT2RP2002618	2813	R-NT2RP2002618	7891
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		NT2RP2002643	F-NT2RP2002643	2815	R-NT2RP2002643	7893
		NT2RP2002672	F-NT2RP2002672	2816	R-NT2RP2002672	7894
	50	NT2RP2002701	F-NT2RP2002701	2817	R-NT2RP2002701	7895
		NT2RP2002706	F-NT2RP2002706	2818	R-NT2RP2002706	7896
		NT2RP2002710	F-NT2RP2002710	2819	R-NT2RP2002710	7897
		NT2RP2002727	F-NT2RP2002727	2820	R-NT2RP2002727	7898
	55	NT2RP2002736	F-NT2RP2002736	2821	R-NT2RP2002736	7899

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	NT2RP2002741	F-NT2RP2002741	2823	R-NT2RP2002741	7901
	NT2RP2002750	F-NT2RP2002750	2824	R-NT2RP2002750	7902
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	NT2RP2002753	F-NT2RP2002753	2826	R-NT2RP2002753	7904
	NT2RP2002769	F-NT2RP2002769	2827	R-NT2RP2002769	7905
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	NT2RP2002857	F-NT2RP2002857	2831	R-NT2RP2002857	7909
	NT2RP2002862	F-NT2RP2002862	2832	R-NT2RP2002862	7910
46	NT2RP2002880	F-NT2RP2002880	2833	R-NT2RP2002880	7911
15	NT2RP2002891	F-NT2RP2002891	2834	R-NT2RP2002891	7912
	NT2RP2002995	F-NT2RP2002925	2835	R-NT2RP2002925	7913
	NT2RP2002928	F-NT2RP2002928	2836	R-NT2RP2002928	7914
			2837	R-NT2RP2002929	7914
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	NT2RP2002954	F-NT2RP2002954	2839	R-NT2RP2002954	
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25	NT2RP2002979	F-NT2RP2002979	2841	R-NT2RP2002979	7918
	NT2RP2002980	F-NT2RP2002980	2842	R-NT2RP2002980	7919
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30	NT2RP2003000	F-NT2RP2003000	2846	R-NT2RP2003000	7923
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	NT2RP2003073	F-NT2RP2003073	2848	R-NT2RP2003073	7925
	NT2RP2003099	F-NT2RP2003099	2849	R-NT2RP2003099	7926
35	NT2RP2003108	F-NT2RP2003108	2850	R-NT2RP2003108	7927
	NT2RP2003117	F-NT2RP2003117	2851	R-NT2RP2003117	7928
	NT2RP2003121	F-NT2RP2003121	2852	R-NT2RP2003121	7929
	NT2RP2003125	F-NT2RP2003125	2853	R-NT2RP2003125	7930
40	NT2RP2003129	F-NT2RP2003129	2854	R-NT2RP2003129	7931
	NT2RP2003137	F-NT2RP2003137	2855	R-NT2RP2003137	7932
	NT2RP2003157	F-NT2RP2003157	2856		
	NT2RP2003158	F-NT2RP2003158	2857		
	NT2RP2003161	F-NT2RP2003161	2858	R-NT2RP2003161	7933
45	NT2RP2003164	F-NT2RP2003164	2859	R-NT2RP2003164	7934
	NT2RP2003165	F-NT2RP2003165	2860	R-NT2RP2003165	7935
	NT2RP2003177	F-NT2RP2003177	2861	R-NT2RP2003177	7936
	NT2RP2003194	F-NT2RP2003194	2862	R-NT2RP2003194	7937
50	NT2RP2003206	F-NT2RP2003206	2863	R-NT2RP2003206	7938
	NT2RP2003228	F-NT2RP2003228	2864		
	NT2RP2003230	F-NT2RP2003230	2865	R-NT2RP2003230	7939
	NT2RP2003237	F-NT2RP2003237	2866	R-NT2RP2003237	7940
55	NT2RP2003243	F-NT2RP2003243	2867	R-NT2RP2003243	7941
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	NT2RP2003272	F-NT2RP2003272	2869	R-NT2RP2003272	7943
	NT2RP2003277	F-NT2RP2003277	2870	R-NT2RP2003277	7944
5	NT2RP2003280	F-NT2RP2003280	2871	R-NT2RP2003280	7945
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	NT2RP2003293	F-NT2RP2003293	2873	R-NT2RP2003293	7947
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	NT2RP2003339	F-NT2RP2003339	2879	R-NT2RP2003339	7952
15	NT2RP2003347	F-NT2RP2003347	2880	R-NT2RP2003339	7953
	NT2RP2003367	F-NT2RP2003367	2881	R-NT2RP2003347	7954
	NT2RP2003391	F-NT2RP2003391	2882	R-NT2RP2003391	7955
	NT2RP2003391	F-NT2RP2003393	2883	R-NT2RP2003391	
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	NT2RP2003401	F-NT2RP2003401	2885	R-NT2RP2003401	7958
	NT2RP2003433	F-NT2RP2003433	2886	R-NT2RP2003433	7959
	NT2RP2003445	F-NT2RP2003445	2887	R-NT2RP2003445	7960
25	NT2RP2003446	F-NT2RP2003446	2888	R-NT2RP2003446	7961
	NT2RP2003456	F-NT2RP2003456	2889	R-NT2RP2003456	7962
	NT2RP2003466	F-NT2RP2003466	2890	K W12M 2000400	1302
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30	NT2RP2003499	F-NT2RP2003499	2892	R-NT2RP2003499	7964
	NT2RP2003506	F-NT2RP2003506	2893	R-NT2RP2003506	7965
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	NT2RP2003513	F-NT2RP2003513	2895	R-NT2RP2003513	7967
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	NT2RP2003533	F-NT2RP2003533	2898	R-NT2RP2003533	7970
	NT2RP2003543	F-NT2RP2003543	2899	R-NT2RP2003543	7971
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40	NT2RP2003564	F-NT2RP2003564	2901	R-NT2RP2003564	7973
	NT2RP2003567	F-NT2RP2003567	2902		
	NT2RP2003581	F-NT2RP2003581	2903	R-NT2RP2003581	7974
	NT2RP2003596	F-NT2RP2003596	2904	R-NT2RP2003596	7975
45	NT2RP2003604	F-NT2RP2003604	2905	R-NT2RP2003604	7976
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	NT2RP2003643	F-NT2RP2003643	2907	R-NT2RP2003643	7978
	NT2RP2003668	F-NT2RP2003668	2908	R-NT2RP2003668	7979
50	NT2RP2003687	F-NT2RP2003687	2909	R-NT2RP2003687	7980
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	NT2RP2003704	F-NT2RP2003704	2912	R-NT2RP2003704	7983
55	NT2RP2003706	F-NT2RP2003706	2913	R-NT2RP2003706	7984

	NT2RP2003713	F-NT2RP2003713	2914	R-NT2RP2003713	7985
	NT2RP2003714	F-NT2RP2003714	2915	R-NT2RP2003714	7986
	NT2RP2003727	F-NT2RP2003727	2916	R-nnnnnnnnnnn	7987
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	NT2RP2003751	F-NT2RP2003751	2918	R-NT2RP2003751	7989
	NT2RP2003760	F-NT2RP2003760	2919	R-NT2RP2003760	7990
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	NT2RP2003777	F-NT2RP2003777	2923	R-NT2RP2003777	7994
	NT2RP2003781	F-NT2RP2003781	2924	R-NT2RP2003781	7995
46	NT2RP2003793	F-NT2RP2003793	2925	R-NT2RP2003793	7996
15	NT2RP2003825	F-NT2RP2003825	2926	K 111210 2000100	1330
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	NT2RP2003885	F-NT2RP2003885	2931	R-NT2RP2003885	8001
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	NT2RP2003952	F-NT2RP2003952	2933	R-NT2RP2003952	8003
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	NT2RP2003981	F-NT2RP2003981	2936	R-NT2RP2003981	8006
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	NT2RP2003988	F-NT2RP2003988	2939	R-NT2RP2003988	8009
	NT2RP2004013	F-NT2RP2004013	2940		
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35	NT2RP2004041	F-NT2RP2004041	2942	R-NT2RP2004041	8011
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40	NT2RP2004098	F-NT2RP2004098	2946	R-NT2RP2004098	8015
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	NT2RP2004142	F-NT2RP2004142	2948	R-NT2RP2004142	8017
	NT2RP2004152	F-NT2RP2004152	2949	R-NT2RP2004152	8018
	NT2RP2004165	F-NT2RP2004165	2950	R-NT2RP2004165	8019
45	NT2RP2004170	F-NT2RP2004170	2951	R-NT2RP2004170	8020
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	NT2RP2004226	F-NT2RP2004226	2957	R-NT2RP2004226	8026
	NT2RP2004232	F-NT2RP2004232	2958	R-NT2RP2004232	8027
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	NT2RP2004240	F-NT2RP2004240	2960	R-NT2RP2004240	8029
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13	NT2RP2004373	F-NT2RP2004373	2972	R-NT2RP2004373	8041
	NT2RP2004389	F-NT2RP2004389	2973	R-NT2RP2004389	8042
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	NT2RP2004412	F-NT2RP2004412	2978	R-NT2RP2004412	8047
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25	NT2RP2004463	F-NT2RP2004463	2980		
	NT2RP2004476	F-NT2RP2004476	2981	R-NT2RP2004476	8049
	NT2RP2004490	F-NT2RP2004490	2982	R-NT2RP2004490	8050
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	NT2RP2004538	F-NT2RP2004538	2985	R-NT2RP2004538	8053
	NT2RP2004551	F-NT2RP2004551	2986	R-NT2RP2004551	8054
	NT2RP2004568	F-NT2RP2004568	2987	R-NT2RP2004568	8055
35	NT2RP2004580	F-NT2RP2004580	2988	R-NT2RP2004580	8056
••	NT2RP2004587	F-NT2RP2004587	2989	R-NT2RP2004587	8057
	NT2RP2004594	F-NT2RP2004594	2990	R-NT2RP2004594	8058
	NT2RP2004600	F-NT2RP2004600	2991	R-NT2RP2004600	8059
40	NT2RP2004602	F-NT2RP2004602	2992	R-NT2RP2004602	8060
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	NT2RP2004664	F-NT2RP2004664	2995	R-NT2RP2004664	8063
	NT2RP2004675	F-NT2RP2004675	2996	R-NT2RP2004675	8064
45	NT2RP2004681	F-NT2RP2004681	2997	R-NT2RP2004681	8065
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	NT2RP2004710	F-NT2RP2004710	3000	R-NT2RP2004710	8068
50	NT2RP2004736	F-NT2RP2004736	3001	R-NT2RP2004736	8069
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	NT2RP2004767	F-NT2RP2004767	3003	R-NT2RP2004767	8071
	NT2RP2004768	F-NT2RP2004768	3004		
55	NT2RP2004775	F-NT2RP2004775	3005	R-NT2RP2004775	8072

	NT2RP2004791	F-NT2RP2004791	3006	R-NT2RP2004791	8073
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	NT2RP2004861	F-NT2RP2004861	3010		
	NT2RP2004897	F-NT2RP2004897	3011	R-NT2RP2004861	8078
40	NT2RP2004933	F-NT2RP2004933		R-NT2RP2004897	8079
10	NT2RP2004936	F-NT2RP2004936	3013	D NECODDOO 400C	0000
			3014	R-NT2RP2004936	8080
	NT2RP2004959 NT2RP2004961	F-NT2RP2004959 F-NT2RP2004961	3015	R-nnnnnnnnnnnn	8081
			3016	R-NT2RP2004961	8082
15	NT2RP2004962	F-NT2RP2004962	3017	R-NT2RP2004962	8083
	NT2RP2004967	F-NT2RP2004967	3018	R-NT2RP2004967	8084
	NT2RP2004978	F-NT2RP2004978	3019	R-NT2RP2004978	8085
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20	NT2RP2004985	F-NT2RP2004985	3021	R-NT2RP2004985	8087
	NT2RP2004999	F-NT2RP2004999	3022	R-NT2RP2004999	8088
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	NT2RP2005001	F-NT2RP2005001	3024	R-NT2RP2005001	8090
25	NT2RP2005003	F-NT2RP2005003	3025	R-NT2RP2005003	8091
20	NT2RP2005012	F-NT2RP2005012	3026	R-nnnnnnnnnnn	8092
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	NT2RP2005020	F-NT2RP2005020	3028	R-NT2RP2005020	8094
	NT2RP2005022	F-NT2RP2005022	3029		
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	NT2RP2005108	F-NT2RP2005108	3033	R-NT2RP2005108	8098
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	NT2RP2005126 NT2RP2005139	F-NT2RP2005126 F-NT2RP2005139	3035	R-NT2RP2005126	8100
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40	NT2RP2005147	F-NT2RP2005147	3038	R-NT2RP2005144	8103
	NT2RP2005159	F-NT2RP2005159	3039	R-NT2RP2005147	8104
	NT2RP2005162	F-NT2RP2005169	3040	R-NT2RP2005159	8105
	NT2RP2005168	F-NT2RP2005162	3041	R-NT2RP2005162	8106
45	NT2RP2005108	F-NT2RP2005204	3042	R-NT2RP2005168	8107
.•	NT2RP2005204 NT2RP2005227	F-NT2RP2005204 F-NT2RP2005227	3043	R-NT2RP2005204	8108
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	NT2RP2005289	F-NT2RP2005288	3050	R-NT2RP2005288	8115
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	NT2RP2005315	F-NT2RP2005315	3053	R-NT2RP2005315	8118
	NT2RP2005325	F-NT2RP2005325	3054	R-NT2RP2005325	8119
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	NT2RP2005344	F-NT2RP2005344	3056	R-NT2RP2005344	8121
	NT2RP2005354	F-NT2RP2005354	3057	R-NT2RP2005354	8122
	NT2RP2005358	F-NT2RP2005358	3058		
10	NT2RP2005360	F-NT2RP2005360	3059	R-NT2RP2005360	8123
	NT2RP2005393	F-NT2RP2005393	3060	R-NT2RP2005393	8124
	NT2RP2005407	F-NT2RP2005407	3061	R-NT2RP2005407	8125
	NT2RP2005436	F-NT2RP2005436	3062	R-NT2RP2005436	8126
45	NT2RP2005441	F-NT2RP2005441	3063	R-NT2RP2005441	8127
15	NT2RP2005453	F-NT2RP2005453	3064	R-NT2RP2005453	8128
	NT2RP2005457	F-NT2RP2005457	3065	R-NT2RP2005457	8129
	NT2RP2005464	F-NT2RP2005464	3066	R-NT2RP2005464	8130
	NT2RP2005465	F-NT2RP2005465	3067	R-NT2RP2005465	8131
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	NT2RP2005476	F-NT2RP2005476	3069	R-NT2RP2005476	8133
	NT2RP2005490	F-NT2RP2005490	3070	R-NT2RP2005490	8134
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25	NT2RP2005495	F-NT2RP2005495	3072	R-NT2RP2005495	8136
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30	NT2RP2005509	F-NT2RP2005509	3076	R-NT2RP2005509	8140
	NT2RP2005520	F-NT2RP2005520	3077	R-NT2RP2005520	8141
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35	NT2RP2005539	F-NT2RP2005539	3080	R-NT2RP2005539	8144
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	NT2RP2005549	F-NT2RP2005549	3082	R-NT2RP2005549	8146
	NT2RP2005555	F-NT2RP2005555	3083	R-NT2RP2005555	8147
	NT2RP2005557	F-NT2RP2005557	3084	R-NT2RP2005557	8148
40	NT2RP2005581	F-NT2RP2005581	3085	R-NT2RP2005581	8149
	NT2RP2005600	F-NT2RP2005600	3086	R-NT2RP2005600	8150
	NT2RP2005605	F-NT2RP2005605	3087	R-NT2RP2005605	8151
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45	NT2RP2005622	F-NT2RP2005622	3089	R-NT2RP2005622	8153
	NT2RP2005635	F-NT2RP2005635	3090		
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50	NT2RP2005645	F-NT2RP2005645	3093	R-NT2RP2005645	8156
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	NT2RP2005654	F-NT2RP2005654	3095	R-NT2RP2005654	8158
	NT2RP2005669	F-NT2RP2005669	3096	R-NT2RP2005669	8159
55	NT2RP2005675	F-NT2RP2005675	3097	R-NT2RP2005675	8160

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	NT2RP2005694	F-NT2RP2005694	3100	R-NT2RP2005694	8163
5	NT2RP2005701	F-NT2RP2005701	3101	R-NT2RP2005701	8164
	NT2RP2005712	F-NT2RP2005712	3102	R-NT2RP2005712	8165
	NT2RP2005719	F-NT2RP2005719	3103	R-NT2RP2005719	8166
	NT2RP2005722	F-NT2RP2005722	3104	R-NT2RP2005722	8167
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1 5	NT2RP3000919	F-NT2RP3000919	3273	R-NT2RP3000919	8327
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50	NT2RP3001004	F-NT2RP3001004	3277	R-NT2RP3001004	8331
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	NT2RP3001339	F-NT2RP3001339	3317	R-NT2RP3001339	8370
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	NT2RP3001428	F-NT2RP3001428	3332	R-nnnnnnnnnnn	8384
	NT2RP3001432	F-NT2RP3001432	3333	R-NT2RP3001432	8385
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	NT2RP3001457	F-NT2RP3001457	3337	R-NT2RP3001457	8389
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55	171214 0003040	1 1112/0 3003040	5551	A HILLIU OVOOTO	5555

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	NT2RP3004480	F-NT2RP3004480	3604	R-NT2RP3004480	8639
	NT2RP3004490	F-NT2RP3004490	3605	R-NT2RP3004490	8640
	NT2RP3004498	F-NT2RP3004498	3606	R-NT2RP3004498	8641
5	NT2RP3004503	F-NT2RP3004503	3607	R-NT2RP3004503	8642
	NT2RP3004504	F-NT2RP3004504	3608	R-NT2RP3004504	8643
	NT2RP3004507	F-NT2RP3004507	3609	R-NT2RP3004507	8644
	NT2RP3004527	F-NT2RP3004527	3610	R-NT2RP3004527	8645
10	NT2RP3004534	F-NT2RP3004534	3611	R-nnnnnnnnnnn	8646
	NT2RP3004539	F-NT2RP3004539	3612		
	NT2RP3004544	F-NT2RP3004544	3613	R-NT2RP3004544	8647
	NT2RP3004566	F-NT2RP3004566	3614	R-NT2RP3004566	8648
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	NT2RP3004572	F-NT2RP3004572	3616	R-NT2RP3004572	8650
	NT2RP3004578	F-NT2RP3004578,	3617	R-NT2RP3004578	8651
	NT2RP3004594	F-NT2RP3004594	3618	R-NT2RP3004594	8652
20	NT2RP3004617	F-NT2RP3004617	3619	R-NT2RP3004617	8653
20	NT2RP3004618	F-NT2RP3004618	3620	R-NT2RP3004618	8654
	NT2RP3004669	F-NT2RP3004669	3621		
	NT2RP3004670	F-NT2RP3004670	3622	R-NT2RP3004670	8655
05	NT2RP4000008	F-NT2RP4000008	3623	R-NT2RP4000008	8656
25	NT2RP4000023	F-NT2RP4000023	3624	R-NT2RP4000023	8657
	NT2RP4000035	F-NT2RP4000035	3625	R-NT2RP4000035	8658
	NT2RP4000049	F-NT2RP4000049	3626	R-NT2RP4000049	8659
	NT2RP4000051	F-NT2RP4000051	3627	R-NT2RP4000051	8660
30	NT2RP4000078	F-NT2RP4000078	3628	R-NT2RP4000078	8661
	NT2RP4000102	F-NT2RP4000102	3629	R-NT2RP4000102	8662
	NT2RP4000109	F-NT2RP4000109	3630	R-NT2RP4000109	8663
	NT2RP4000111	F-NT2RP4000111	3631		
35	NT2RP4000129	F-NT2RP4000129	3632	R-NT2RP4000129	8664
	NT2RP4000147	F-NT2RP4000147	3633	R-NT2RP4000147	8665
	NT2RP4000150	F-NT2RP4000150	3634	R-NT2RP4000150	8666
	NT2RP4000151	F-NT2RP4000151	3635	R-NT2RP4000151	8667
40	NT2RP4000159	F-NT2RP4000159	3636	R-NT2RP4000159	8668
	NT2RP4000167	F-NT2RP4000167	3637	R-NT2RP4000167	8669
	NT2RP4000185	F-NT2RP4000185	3638	R-NT2RP4000185	8670
	NT2RP4000210	F-NT2RP4000210	3639	R-NT2RP4000210	8671
45	NT2RP4000212	F-NT2RP4000212	3640	R-NT2RP4000212	8672
	NT2RP4000214	F-NT2RP4000214	3641	R-NT2RP4000214	8673
	NT2RP4000218	F-NT2RP4000218	3642	R-NT2RP4000218	8674
	NT2RP4000243	F-NT2RP4000243	3643	R-NT2RP4000243	8675
50	NT2RP4000246	F-NT2RP4000246	3644	R-NT2RP4000246	8676
	NT2RP4000259	F-NT2RP4000259	3645	R-NT2RP4000259	8677
	NT2RP4000263	F-NT2RP4000263	3646	R-NT2RP4000263	8678
	NT2RP4000290	F-NT2RP4000290	3647	R-nnnnnnnnnnnn	8679
	NT2RP4000312 NT2RP4000321	F-NT2RP4000312	3648	R-NT2RP4000312	8680
55	N12KP4000321	F-NT2RP4000321	3649	R-NT2RP4000321	8681

	NT2RP4000323	F-NT2RP4000323	3650	R-NT2RP4000323	8682
	NT2RP4000355	F-NT2RP4000355	3651	R-NT2RP4000355	8683
	NT2RP4000360	F-NT2RP4000360	3652	R-NT2RP4000360	8684
5	NT2RP4000367	F-NT2RP4000367	3653	R-NT2RP4000367	8685
	NT2RP4000370	F-NT2RP4000370	3654	R-NT2RP4000370	8686
	NT2RP4000376	F-NT2RP4000376	3655	R-NT2RP4000376	8687
	NT2RP4000381	F-NT2RP4000381	3656	R-NT2RP4000381	8688
10	NT2RP4000398	F-NT2RP4000398	3657		
	NT2RP4000415	F-NT2RP4000415	3658	R-NT2RP4000415	8689
	NT2RP4000417	F-NT2RP4000417	3659	R-NT2RP4000417	8690
	NT2RP4000424	F-NT2RP4000424	3660	R-NT2RP4000424	8691
15	NT2RP4000448	F-NT2RP4000448	3661	R-NT2RP4000448	8692
15	NT2RP4000449	F-NT2RP4000449	3662	R-NT2RP4000449	8693
	NT2RP4000455	F-NT2RP4000455,	3663	R-NT2RP4000455	8694
	NT2RP4000457	F-NT2RP4000457	3664	R-nnnnnnnnnnn	8695
	NT2RP4000480	F-NT2RP4000480	3665	R-NT2RP4000480	8696
20	NT2RP4000481	F-NT2RP4000481	3666	R-nnnnnnnnnnn	8697
	NT2RP4000498	F-NT2RP4000498	3667	к иншишиши	0051
	NT2RP4000500	F-NT2RP4000500	3668	R-NT2RP4000500	8698
	NT2RP4000515	F-NT2RP4000515	3669	R-NT2RP4000515	8699
25	NT2RP4000517	F-NT2RP4000517	3670	R-NT2RP4000517	8700
	NT2RP4000518	F-NT2RP4000518	3671	R-NT2RP4000518	8701
	NT2RP4000519	F-NT2RP4000519	3672	R-NT2RP4000519	8702
	NT2RP4000524	F-NT2RP4000524	3673	R-NT2RP4000524	8703
30	NT2RP4000528	F-NT2RP4000528	3674	R-NT2RP4000528	8704
	NT2RP4000541	F-NT2RP4000541	3675	R-NT2RP4000541	8705
	NT2RP4000556	F-NT2RP4000556	3676	R-NT2RP4000556	8706
	NT2RP4000560	F-NT2RP4000560	3677		
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	NT2RP4000614	F-NT2RP4000614	3679	R-NT2RP4000614	8708
	NT2RP4000638	F-NT2RP4000638	3680	R-NT2RP4000638	8709
	NT2RP4000648	F-NT2RP4000648	3681	R-NT2RP4000648	8710
40	NT2RP4000657	F-NT2RP4000657	3682	R-NT2RP4000657	8711
10	NT2RP4000704	F-NT2RP4000704	3683	R-NT2RP4000704	8712
	NT2RP4000713	F-NT2RP4000713	3684		
	NT2RP4000724	F-NT2RP4000724	3685	R-NT2RP4000724	8713
	NT2RP4000728	F-NT2RP4000728	3686	R-NT2RP4000728	8714
45	NT2RP4000737	F-NT2RP4000737	3687		
	NT2RP4000739	F-NT2RP4000739	3688	R-NT2RP4000739	8715
	NT2RP4000781	F-NT2RP4000781	3689	R-NT2RP4000781	8716
	NT2RP4000787	F-NT2RP4000787	3690		
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	NT2RP4000833	F-NT2RP4000833	3692	R-NT2RP4000833	8718
	NT2RP4000837	F-NT2RP4000837	3693	R-NT2RP4000837	8719
	NT2RP4000839	F-NT2RP4000839	3694		
55	NT2RP4000855	F-NT2RP4000855	3695	R-NT2RP4000855	8720

	NT2RP4000865	F-NT2RP4000865	3696	R-NT2RP4000865	8721
	NT2RP4000878	F-NT2RP4000878	3697	R-NT2RP4000878	8722
	NT2RP4000879	F-NT2RP4000879	3698	R-NT2RP4000879	8723
5	NT2RP4000907	F-NT2RP4000907	3699	R-nnnnnnnnnnn	8724
	NT2RP4000915	F-NT2RP4000915	3700	R-nnnnnnnnnnn	8725
	NT2RP4000918	F-NT2RP4000918	3701		
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10	NT2RP4000927	F-NT2RP4000927	3703	R-nnnnnnnnnnn	8727
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	NT2RP4000929	F-NT2RP4000929	3705	R-NT2RP4000929	8729
	NT2RP4000955	F-NT2RP4000955	3706	R-NT2RP4000955	8730
15	NT2RP4000973	F-NT2RP4000973	3707	R-NT2RP4000973	8731
	NT2RP4000975	F-NT2RP4000975	3708	R-NT2RP4000975	8732
	NT2RP4000979	F-NT2RP4000979,	3709	R-NT2RP4000979	8733
	NT2RP4000984	F-NT2RP4000984	3710	R-NT2RP4000984	8734
20	NT2RP4000989	F-NT2RP4000989	3711	R-NT2RP4000989	8735
20	NT2RP4000996	F-NT2RP4000996	3712	R-NT2RP4000996	8736
	NT2RP4000997	F-NT2RP4000997	3713	R-NT2RP4000997	8737
	NT2RP4001004	F-NT2RP4001004	3714	R-NT2RP4001004	8738
	NT2RP4001006	F-NT2RP4001006	3715	R-NT2RP4001006	8739
25	NT2RP4001010	F-NT2RP4001010	3716	R-NT2RP4001010	8740
	NT2RP4001029	F-NT2RP4001029	3717	R-NT2RP4001029	8741
	NT2RP4001041	F-NT2RP4001041	3718	R-NT2RP4001041	8742
	NT2RP4001057	F-NT2RP4001057	3719	R-NT2RP4001057	8743
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	NT2RP4001078	F-NT2RP4001078	3721	R-NT2RP4001078	8745
	NT2RP4001079	F-NT2RP4001079	3722	R-NT2RP4001079	8746
	NT2RP4001080	F-NT2RP4001080	3723	R-NT2RP4001080	8747
35	NT2RP4001086	F-NT2RP4001086	3724	R-nnnnnnnnnnn	8748
	NT2RP4001095	F-NT2RP4001095	3725	R-NT2RP4001095	8749
	NT2RP4001100	F-NT2RP4001100	3726	R-NT2RP4001100	8750
	NT2RP4001117	F-NT2RP4001117	3727	R-NT2RP4001117	8751
40	NT2RP4001122	F-NT2RP4001122	3728	R-NT2RP4001122	8752
	NT2RP4001126	F-NT2RP4001126	3729	R-NT2RP4001126	8753
	NT2RP4001138	F-NT2RP4001138	3730	R-NT2RP4001138	8754
	NT2RP4001143	F-NT2RP4001143	3731	R-NT2RP4001143	8755
45	NT2RP4001148	F-NT2RP4001148	3732	R-NT2RP4001148	8756
	NT2RP4001149	F-NT2RP4001149	3733	R-NT2RP4001149	8757
	NT2RP4001150	F-NT2RP4001150	3734	R-NT2RP4001150	8758
	NT2RP4001159	F-NT2RP4001159	3735	R-NT2RP4001159	8759
50	NT2RP4001174	F-NT2RP4001174	3736	R-NT2RP4001174	8760
50	NT2RP4001206	F-NT2RP4001206	3737	R-nnnnnnnnnnn	8761
	NT2RP4001207	F-NT2RP4001207	3738	R-NT2RP4001207	8762
	NT2RP4001210	F-NT2RP4001210	3739	R-NT2RP4001210	8763
	NT2RP4001213	F-NT2RP4001213	3740	R-NT2RP4001213	8764
55	NT2RP4001219	F-NT2RP4001219	3741	R-NT2RP4001219	8765

	NT2RP4001228	F-NT2RP4001228	3742	R-NT2RP4001228	8766
	NT2RP4001235	F-NT2RP4001235	3743	R-NT2RP4001235	8767
	NT2RP4001256	F-NT2RP4001256	3744	R-NT2RP4001256	8768
5	NT2RP4001260	F-NT2RP4001260	3745	R-NT2RP4001260	8769
	NT2RP4001274	F-NT2RP4001274	3746	R-NT2RP4001274	8770
	NT2RP4001276	F-NT2RP4001276	3747	R-nnnnnnnnnnn	8771
	NT2RP4001313	F-NT2RP4001313	3748	R-NT2RP4001313	8772
10	NT2RP4001315	F-NT2RP4001315	3749	R-NT2RP4001315	8773
	NT2RP4001336	F-NT2RP4001336	3750		
	NT2RP4001339	F-NT2RP4001339	3751	R-NT2RP4001339	8774
	NT2RP4001343	F-NT2RP4001343	3752		
15	NT2RP4001345	F-NT2RP4001345	3753	R-NT2RP4001345	8775
	NT2RP4001351	F-NT2RP4001351	3754	R-NT2RP4001351	8776
	NT2RP4001353	F-NT2RP4001353,	3755	R-NT2RP4001353	8777
	NT2RP4001372	F-NT2RP4001372	3756	R-NT2RP4001372	8778
20	NT2RP4001373	F-NT2RP4001373	3757	R-NT2RP4001373	8779
20	NT2RP4001375	F-NT2RP4001375	3758	R-NT2RP4001375	8780
	NT2RP4001379	F-NT2RP4001379	3759	R-NT2RP4001379	8781
	NT2RP4001389	F-NT2RP4001389	3760	R-NT2RP4001389	8782
	NT2RP4001407	F-NT2RP4001407	3761	R-NT2RP4001407	8783
25	NT2RP4001414	F-NT2RP4001414	3762	R-NT2RP4001414	8784
	NT2RP4001433	F-NT2RP4001433	3763	R-NT2RP4001433	8785
	NT2RP4001442	F-NT2RP4001442	3764	R-NT2RP4001442	8786
	NT2RP4001447	F-NT2RP4001447	3765	R-NT2RP4001447	8787
30	NT2RP4001474	F-NT2RP4001474	3766	R-NT2RP4001474	8788
	NT2RP4001483	F-NT2RP4001483	3767	R-NT2RP4001483	8789
	NT2RP4001498	F-NT2RP4001498	3768	R-NT2RP4001498	8790
	NT2RP4001502	F-NT2RP4001502	3769	R-NT2RP4001502	8791
35	NT2RP4001507	F-NT2RP4001507	3770	R-NT2RP4001507	8792
	NT2RP4001524	F-NT2RP4001524	3771	R-NT2RP4001524	8793
	NT2RP4001529	F-NT2RP4001529	3772	R-NT2RP4001529	8794
	NT2RP4001547	F-NT2RP4001547	3773	R-NT2RP4001547	8795
40	NT2RP4001551	F-NT2RP4001551	3774	R-nnnnnnnnnnn	8796
	NT2RP4001555	F-NT2RP4001555	3775	R-NT2RP4001555	8797
	NT2RP4001567	F-NT2RP4001567	3776	R-NT2RP4001567	8798
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45	NT2RP4001571	F-NT2RP4001571	3778	R-NT2RP4001571	8800
40	NT2RP4001574	F-NT2RP4001574	3779	R-NT2RP4001574	8801
	NT2RP4001575	F-NT2RP4001575	3780	R-NT2RP4001575	8802
	NT2RP4001592	F-NT2RP4001592	3781	R-NT2RP4001592	8803
50	NT2RP4001610	F-NT2RP4001610	3782	R-NT2RP4001610	8804
50	NT2RP4001614	F-NT2RP4001614	3783	R-NT2RP4001614	8805
	NT2RP4001634	F-NT2RP4001634	3784	R-NT2RP4001634	8806
	NT2RP4001638	F-NT2RP4001638	3785	R-NT2RP4001638	8807
	NT2RP4001644	F-NT2RP4001644	3786	R-NT2RP4001644	8808
55	NT2RP4001656	F-NT2RP4001656	3787	R-NT2RP4001656	8809

	NT2RP4001677	F-NT2RP4001677	3788	R-NT2RP4001677	8810
	NT2RP4001679	F-NT2RP4001679	3789		
	NT2RP4001696	F-NT2RP4001696	3790	R-NT2RP4001696	8811
5	NT2RP4001725	F-NT2RP4001725	3791	R-NT2RP4001725	8812
	NT2RP4001730	F-NT2RP4001730	3792	R-nnnnnnnnnnn	8813
	NT2RP4001739	F-NT2RP4001739	3793	R-NT2RP4001739	8814
	NT2RP4001753	F-NT2RP4001753	3794	R-NT2RP4001753	8815
10	NT2RP4001760	F-NT2RP4001760	3795	R-NT2RP4001760	8816
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	NT2RP4001803	F-NT2RP4001803	3797	R-NT2RP4001803	8818
	NT2RP4001822	F-NT2RP4001822	3798	R-NT2RP4001822	8819
15	NT2RP4001823	F-NT2RP4001823	3799	R-NT2RP4001823	8820
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	NT2RP4001838	F-NT2RP4001838.	3801	R-NT2RP4001838	8822
	NT2RP4001841	F-NT2RP4001841	3802		
	NT2RP4001849	F-NT2RP4001849	3803	R-NT2RP4001849	8823
20	NT2RP4001861	F-NT2RP4001861	3804		
	NT2RP4001889	F-NT2RP4001889	3805	R-NT2RP4001889	8824
	NT2RP4001893	F-NT2RP4001893	3806	R-NT2RP4001893	8825
	NT2RP4001896	F-NT2RP4001896	3807	R-NT2RP4001896	8826
25	NT2RP4001901	F-NT2RP4001901	3808	R-NT2RP4001901	8827
	NT2RP4001927	F-NT2RP4001927	3809	R-NT2RP4001927	8828
	NT2RP4001938	F-NT2RP4001938	3810	R-NT2RP4001938	8829
	NT2RP4001946	F-NT2RP4001946	3811	R-NT2RP4001946	8830
30	NT2RP4001950	F-NT2RP4001950	3812	R-NT2RP4001950	8831
	NT2RP4001953	F-NT2RP4001953	3813	R-NT2RP4001953	8832
	NT2RP4001966	F-NT2RP4001966	3814	R-NT2RP4001966	8833
	NT2RP4001975	F-NT2RP4001975	3815	R-NT2RP4001975	8834
35	NT2RP4002018	F-NT2RP4002018	3816	R-NT2RP4002018	8835
	NT2RP4002047	F-NT2RP4002047	3817	R-NT2RP4002047	8836
	NT2RP4002052	F-NT2RP4002052	3818	R-NT2RP4002052	8837
	NT2RP4002058	F-NT2RP4002058	3819	R-NT2RP4002058	8838
40	NT2RP4002071	F-NT2RP4002071	3820	R-NT2RP4002071	8839
	NT2RP4002075	F-NT2RP4002075	3821	R-NT2RP4002075	8840
	NT2RP4002078	F-NT2RP4002078	3822	R-NT2RP4002078	8841
	NT2RP4002081	F-NT2RP4002081	3823	R-nnnnnnnnnnn	8842
45	NT2RP4002083	F-NT2RP4002083	3824	R-NT2RP4002083	8843
40	NT2RP4002408	F-NT2RP4002408	3825	R-NT2RP4002408	8844
	NT2RP4002791	F-NT2RP4002791	3826	R-NT2RP4002791	8845
	NT2RP4002888	F-NT2RP4002888	3827	R-NT2RP4002888	8846
	NT2RP4002905	F-NT2RP4002905	3828	R-NT2RP4002905	8847
50	NT2RP5003459	F-NT2RP5003459	3829		
	NT2RP5003461	F-NT2RP5003461	3830		
	NT2RP5003477	F-NT2RP5003477	3831		
	NT2RP5003492	F-NT2RP5003492	3832		
55	NT2RP5003500	F-NT2RP5003500	3833		

	NT2RP5003506	F-NT2RP5003506	3834		
	NT2RP5003512	F-NT2RP5003512	3835		
	NT2RP5003522	F-NT2RP5003522	3836		
5	NT2RP5003524	F-NT2RP5003524	3837		
	NT2RP5003534	F-NT2RP5003534	3838		
	0VARC1000001	F-0VARC1000001	3839	R-0VARC1000001	8848
	0VARC1000004	F-0VARC1000004	3840	R-0VARC1000004	8849
10	OVARC1000006	F-0VARC1000006	3841	R-0VARC1000006	8850
	0VARC1000013	F-0VARC1000013	3842	R-0VARC1000013	8851
	OVARC1000014	F-0VARC1000014	3843	R-OVARC1000014	8852
	OVARC1000017	F-0VARC1000017	3844	R-0VARC1000017	8853
15	0VARC1000035	F-0VARC1000035	3845	R-0VARC1000035	8854
	OVARC1000058	F-0VARC1000058	3846	R-OVARC1000058	8855
	0VARC1000060	F-0VARC1000060,	3847	R-0VARC1000060	8856
	0VARC1000068	F-0VARC1000068	3848	R-0VARC1000068	8857
20	OVARC1000071	F-0VARC1000071	3849	R-OVARC1000071	8858
20	0VARC1000085	F-0VARC1000085	3850	R-OVARC1000085	8859
	OVARC1000087	F-0VARC1000087	3851	R-nnnnnnnnnnn	8860
	OVARC1000091	F-0VARC1000091	3852	R-0VARC1000091	8861
	OVARC1000092	F-0VARC1000092	3853	R-OVARC1000092	8862
25	OVARC1000106	F-0VARC1000106	3854	R-OVARC1000106	8863
	OVARC1000109	F-0VARC1000109	3855		
	OVARC1000113	F-0VARC1000113	3856	R-OVARC1000113	8864
	OVARC1000114	F-0VARC1000114	3857	R-0VARC1000114	8865
30	OVARC1000133	F-0VARC1000133	3858	R-0VARC1000133	8866
	OVARC1000139	F-0VARC1000139	3859		
	0VARC1000145	F-0VARC1000145	3860	R-OVARC1000145	8867
	0VARC1000148	F-0VARC1000148	3861	R-0VARC1000148	8868
35	OVARC1000151	F-0VARC1000151	3862	R-0VARC1000151	8869
	OVARC1000168	F-0VARC1000168	3863	R-0VARC1000168	8870
	OVARC1000191	F-0VARC1000191	3864	R-OVARC1000191	8871
	OVARC1000198 OVARC1000209	F-0VARC1000198	3865	R-OVARC1000198	8872
40	0VARC1000209 0VARC1000212	F-0VARC1000209 F-0VARC1000212	3866	R-OVARC1000209	8873
	0VARC1000212	F-0VARC1000212	3867 3868	R-OVARC1000212	8874
	0VARC1000240 0VARC1000241	F-0VARC1000240 F-0VARC1000241	3869	R-0VARC1000240	8875
	0VARC1000241	F-0VARC1000241	3870	R-OVARC1000241 R-OVARC1000288	8876
45	0VARC1000288	F-0VARC1000288	3871	R-0VARC1000288	8877 8878
	0VARC1000302	F-0VARC1000302	3872	R-0VARC1000302	8879
	0VARC1000304	F-0VARC1000309	3873	R-0VARC1000304	8880
	OVARC1000303	F-0VARC1000303	3874	R-0VARC1000309	8881
50	0VARC1000326	F-0VARC1000326	3875	R-0VARC1000321	8882
	0VARC1000335	F-0VARC1000325	3876	R-0VARC1000325	8883
	0VARC1000347	F-0VARC1000347	3877	R-OVARC1000333	8884
	0VARC1000384	F-0VARC1000384	3878	R-0VARC1000384	8885
55	0VARC1000408	F-0VARC1000408	3879	R-0VARC1000408	8886
55			55.0	" ANTIOIONATAN	5500

	OVARC1000411	F-0VARC1000411	3880	R-OVARC1000411	8887
	OVARC1000414	F-0VARC1000414	3881	R-0VARC1000414	8888
	OVARC1000420	F-0VARC1000420	3882	R-OVARC1000420	8889
5	0VARC1000427	F-0VARC1000427	3883	R-0VARC1000427	8890
	0VARC1000431	F-0VARC1000431	3884	R-OVARC1000431	8891
	0VARC1000437	F-0VARC1000437	3885	R-0VARC1000437	8892
	OVARC1000440	F-0VARC1000440	3886	R-0VARC1000440	8893
10	0VARC1000442	F-0VARC1000442	3887	R-OVARC1000442	8894
	0VARC1000443	F-0VARC1000443	3888	R-OVARC1000443	8895
	0VARC1000461	F-0VARC1000461	3889	R-OVARC1000461	8896
	0VARC1000465	F-0VARC1000465	3890	R-OVARC1000465	8897
15	0VARC1000466	F-0VARC1000466	3891	R-0VARC1000466	8898
,,	0VARC1000473	F-0VARC1000473	3892	R-0VARC1000473	8899
	0VARC1000479	F-0VARC1000479,	3893	R-OVARC1000479	8900
	0VARC1000486	F-OVARC1000486	3894	R-0VARC1000486	8901
	0VARC1000496	F-0VARC1000496	3895	R-0VARC1000496	8902
20	0VARC1000520	F-0VARC1000520	3896	R-OVARC1000520	8903
	0VARC1000526	F-0VARC1000526	3897	R-0VARC1000526	8904
	0VARC1000533	F-0VARC1000533	3898	R-0VARC1000533	8905
	0VARC1000543	F-0VARC1000543	3899	R-0VARC1000543	8906
25	0VARC1000556	F-0VARC1000556	3900	R-0VARC1000556	8907
	0VARC1000557	F-0VARC1000557	3901	R-OVARC1000557	8908
	0VARC1000564	F-0VARC1000564	3902	R-0VARC1000564	8909
	0VARC1000573	F-0VARC1000573	3903	R-OVARC1000573	8910
30	0VARC1000576	F-0VARC1000576	3904		
	0VARC1000578	F-0VARC1000578	3905	R-0VARC1000578	8911
	0VARC1000588	F-0VARC1000588	3906	R-0VARC1000588	8912
	0VARC1000605	F-0VARC1000605	3907	R-0VARC1000605	8913
35	0VARC1000622	F-0VARC1000622	3908	R-OVARC1000622	8914
	0VARC1000640	F-0VARC1000640	3909	R-0VARC1000640	8915
	0VARC1000649	F-0VARC1000649	3910		
	0VARC1000661	F-0VARC1000661	3911	R-OVARC1000661	8916
40	0VARC1000678	F-0VARC1000678	3912	R-OVARC1000678	8917
	0VARC1000679	F-0VARC1000679	3913	R-nnnnnnnnnnn	8918
	OVARC1000681	F-0VARC1000681	3914	R-OVARC1000681	8919
	0VARC1000682	F-0VARC1000682	3915		
	0VARC1000689	F-0VARC1000689	3916	R-0VARC1000689	8920
45	0VARC1000700	F-0VARC1000700	3917	R-0VARC1000700	8921
	OVARC1000703	F-0VARC1000703	3918	R-0VARC1000703	8922
	OVARC1000722	F-0VARC1000722	3919		
	OVARC1000730	F-0VARC1000730	3920	R-OVARC1000730	8923
50	OVARC1000746	F-0VARC1000746	3921	R-OVARC1000746	8924
	OVARC1000769	F-0VARC1000769	3922	R-0VARC1000769	8925
	0VARC1000771	F-0VARC1000771	3923	R-OVARC1000771	8926
	OVARC1000781	F-0VARC1000781	3924	R-0VARC1000781	8927
55	0VARC1000787	F-0VARC1000787	3925	R-OVARC1000787	8928

	OVARC1000800	F-0VARC1000800	3926	R-0VARC1000800	8929
	OVARC1000802	F-0VARC1000802	3927	R-0VARC1000802	8930
	OVARC1000834	F-0VARC1000834	3928	R-0VARC1000834	8931
5	OVARC1000846	F-0VARC1000846	3929	R-0VARC1000834	
	OVARC1000850	F-0VARC1000850	3930		8932
	OVARC1000862	F-0VARC1000862	3931	R-0VARC1000850	8933
	OVARC1000876	F-0VARC1000802	3932	R-0VARC1000862	8934
10	OVARC1000810	F-0VARC1000870	3933	R-0VARC1000876 R-0VARC1000883	8935
70	OVARC1000885	F-0VARC1000885	3934		8936
	OVARC1000886	F-0VARC1000886	3935	R-0VARC1000885	8937
	OVARC1000890	F-0VARC1000880		R-0VARC1000886	8938
	0VARC1000890		3936	D 0V4DC100001	0000
15	OVARC1000897	F-0VARC1000891	3937	R-0VARC1000891	8939
	OVARC1000897	F-0VARC1000897	3938	R-0VARC1000897	8940
	OVARC1000912	F-OVARC1000912, F-OVARC1000915	3939	R-0VARC1000912	8941
	OVARC1000915 OVARC1000924		3940	R-0VARC1000915	8942
20		F-0VARC1000924	3941	R-0VARC1000924	8943
	OVARC1000936 OVARC1000937	F-0VARC1000936	3942	R-0VARC1000936	8944
	OVARC1000937	F-0VARC1000937	3943	R-0VARC1000937	8945
	0VARC1000948	F-0VARC1000945	3944	R-0VARC1000945	8946
25	0VARC1000948	F-0VARC1000948 F-0VARC1000959	3945	R-0VARC1000948	8947
	OVARC1000959	F-0VARC1000959	3946	R-0VARC1000959	8948
	OVARC1000960 OVARC1000964	F-0VARC1000960 F-0VARC1000964	3947 3948	R-0VARC1000960	8949
	OVARC1000971	F-0VARC1000904	3949	D_0VADC1000071	9050
30	OVARC1000984	F-0VARC1000971	3950	R-0VARC1000971 R-0VARC1000984	8950 8951
	OVARC1000996	F-0VARC1000996	3951	R-0VARC1000984 R-0VARC1000996	8952
	OVARC1000999	F-0VARC1000999	3952	R-0VARC1000998	8953
	OVARC1001000	F-OVARC1001000	3953	R-0VARC1000999	8954
	OVARC1001004	F-0VARC1001004	3954	R-0VARC1001000	8955
35	OVARC1001010	F-0VARC1001010	3955	R-0VARC1001004	8956
	OVARC1001011	F-0VARC1001011	3956	R-0VARC1001011	8957
	OVARC1001032	F-0VARC1001032	3957	R-0VARC1001011	8958
	0VARC1001034	F-0VARC1001034	3958	R-0VARC1001032	8959
40	OVARC1001038	F-0VARC1001038	3959	R-0VARC1001034	8960
	OVARC1001040	F-0VARC1001040	3960	R-0VARC1001040	8961
	0VARC1001044	F-0VARC1001044	3961	R-0VARC1001044	8962
	OVARC1001051	F-0VARC1001051	3962	R-0VARC1001051	8963
45	OVARC1001055	F-0VARC1001055	3963	R-0VARC1001055	8964
	OVARC1001062	F-0VARC1001062	3964	R-0VARC1001062	8965
	OVARC1001065	F-0VARC1001065	3965		0000
	OVARC1001068	F-0VARC1001068	3966	R-0VARC1001068	8966
50	OVARC1001072	F-0VARC1001072	3967	R-OVARC1001072	8967
	OVARC1001074	F-0VARC1001074	3968	R-OVARC1001074	8968
	OVARC1001085	F-0VARC1001085	3969	R-0VARC1001085	8969
	OVARC1001092	F-0VARC1001092	3970	R-0VARC1001092	8970
55	OVARC1001107	F-0VARC1001107	3971		
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	OVARC1001113	F-0VARC1001113	3972	R-0VARC1001113	8971
	OVARC1001117	F-0VARC1001117	3973	R-0VARC1001117	8972
	OVARC1001118	F-0VARC1001118	3974	R-0VARC1001118	8973
5	OVARC1001129	F-0VARC1001129	3975	R-0VARC1001129	8974
	OVARC1001154	F-0VARC1001154	3976		
	OVARC1001161	F-0VARC1001161	3977	R-0VARC1001161	8975
	OVARC1001162	F-0VARC1001162	3978	R-0VARC1001162	8976
10	OVARC1001167	F-0VARC1001167	3979	R-0VARC1001167	8977
	OVARC1001169	F-0VARC1001169	3980	R-0VARC1001169	8978
	OVARC1001170	F-0VARC1001170	3981	R-0VARC1001170	8979
	OVARC1001171	F-0VARC1001171	3982	0114101001110	5510
46	OVARC1001173	F-0VARC1001173	3983	R-0VARC1001173	8980
15	OVARC1001176	F-0VARC1001176	3984	K 07/M01001110	0300
	OVARC1001180	F-OVARC1001180,	3985	R-0VARC1001180	8981
	OVARC1001188	F-0VARC1001188	3986	R-0VARC1001188	8982
	OVARC1001200	F-0VARC1001200	3987	R-0VARC1001200	8983
20	OVARC1001232	F-0VARC1001232	3988	R-OVARC1001232	8984
	OVARC1001240	F-0VARC1001240	3989	R-0VARC1001240	8985
	OVARC1001243	F-0VARC1001243	3990	R-0VARC1001243	8986
	OVARC1001244	F-0VARC1001244	3991	W 0.77001510	0500
25	OVARC1001261	F-0VARC1001261	3992	R-OVARC1001261	8987
	OVARC1001268	F-0VARC1001268	3993	R-0VARC1001268	8988
	OVARC1001270	F-0VARC1001270	3994	R-0VARC1001270	8989
	OVARC1001271	F-0VARC1001271	3995	R-OVARC1001271	8990
30	OVARC1001282	F-0VARC1001282	3996	R-0VARC1001282	8991
	OVARC1001296	F-0VARC1001296	3997	R-0VARC1001296	8992
	OVARC1001306	F-0VARC1001306	3998	R-nnnnnnnnnnn	8993
	OVARC1001329	F-0VARC1001329	3999	R-OVARC1001329	8994
25	OVARC1001330	F-0VARC1001330	4000	R-0VARC1001330	8995
35	OVARC1001339	F-0VARC1001339	4001	R-0VARC1001339	8996
	OVARC1001341	F-0VARC1001341	4002	R-0VARC1001341	8997
	OVARC1001342	F-0VARC1001342	4003	R-OVARC1001342	8998
	OVARC1001344	F-0VARC1001344	4004	R-OVARC1001344	8999
40	OVARC1001357	F-0VARC1001357	4005	R-0VARC1001357	9000
	OVARC1001360	F-0VARC1001360	4006	R-0VARC1001360	9001
	OVARC1001369	F-0VARC1001369	4007	R-OVARC1001369	9002
	OVARC1001372	F-0VARC1001372	4008	R-OVARC1001372	9003
45	OVARC1001376	F-0VARC1001376	4009	R-0VARC1001376	9004
	OVARC1001381	F-0VARC1001381	4010	R-OVARC1001381	9005
	OVARC1001391	F-0VARC1001391	4011	R-0VARC1001391	9006
	OVARC1001399	F-0VARC1001399	4012	R-nnnnnnnnnn	9007
50	OVARC1001417	F-0VARC1001417	4013	R-OVARC1001417	9008
	OVARC1001419	F-0VARC1001419	4014	R-0VARC1001419	9009
	OVARC1001425	F-0VARC1001425	4015	R-0VARC1001425	9010
	OVARC1001436	F-0VARC1001436	4016	R-OVARC1001436	9011
E E	OVARC1001442	F-0VARC1001442	4017	R-0VARC1001442	9012
55					-

	OVARC1001453	F-0VARC1001453	4018	R-0VARC1001453	- 9013
	OVARC1001476	F-0VARC1001476	4019	R-0VARC1001476	9014
	OVARC1001480	F-0VARC1001480	4020	R-0VARC1001480	9015
5	OVARC1001489	F-0VARC1001489	4021	R-0VARC1001489	9016
	OVARC1001496	F-0VARC1001496	4022	R-0VARC1001496	9017
	OVARC1001506	F-0VARC1001506	4023	R-0VARC1001506	9018
	OVARC1001525	F-0VARC1001525	4024	R-0VARC1001525	9019
10	0VARC1001542	F-0VARC1001542	4025	R-0VARC1001542	9020
	OVARC1001547	F-0VARC1001547	4026	R-0VARC1001547	9021
	OVARC1001555	F-0VARC1001555	4027		
	OVARC1001577	F-0VARC1001577	4028	R-OVARC1001577	9022
15	OVARC1001600	F-0VARC1001600	4029	R-0VARC1001600	9023
10	OVARC1001610	F-0VARC1001610	4030	R-0VARC1001610	9024
	OVARC1001611	F-0VARC1001611,	4031	R-0VARC1001611	9025
	OVARC1001615	F-0VARC1001615	4032	R-OVARC1001615	9026
	OVARC1001668	F-0VARC1001668	4033	R-0VARC1001668	9027
20	OVARC1001702	F-0VARC1001702	4034	R-0VARC1001702	9028
	OVARC1001703	F-0VARC1001703	4035	R-OVARC1001703	9029
	OVARC1001711	F-0VARC1001711	4036	R-0VARC1001711	9030
	OVARC1001713	F-0VARC1001713	4037		
25	OVARC1001726	F-0VARC1001726	4038	R-OVARC1001726	9031
	OVARC1001731	F-0VARC1001731	4039	R-OVARC1001731	9032
	OVARC1001745	F-0VARC1001745	4040	R-0VARC1001745	9033
	OVARC1001762	F-0VARC1001762	4041	R-nnnnnnnnnnn	9034
30	OVARC1001766	F-0VARC1001766	4042	R-0VARC1001766	9035
	OVARC1001767	F-0VARC1001767	4043	R-nnnnnnnnnnn	9036
	OVARC1001768	F-0VARC1001768	4044	R-OVARC1001768	9037
	OVARC1001791	F-0VARC1001791	4045	R-OVARC1001791	9038
35	OVARC1001795	F-0VARC1001795	4046	R-0VARC1001795	9039
	OVARC1001802	F-0VARC1001802	4047	R-0VARC1001802	9040
	OVARC1001805	F-0VARC1001805	4048	R-0VARC1001805	9041
	OVARC1001809	F-0VARC1001809	4049		
40	OVARC1001812	F-0VARC1001812	4050	R-OVARC1001812	9042
40	OVARC1001813	F-0VARC1001813	4051	R-OVARC1001813	9043
	OVARC1001820	F-0VARC1001820	4052	R-OVARC1001820	9044
	OVARC1001828	F-0VARC1001828	4053	R-OVARC1001828	9045
	OVARC1001846	F-0VARC1001846	4054	R-OVARC1001846	9046
45	OVARC1001861	F-0VARC1001861	4055	R-0VARC1001861	9047
	OVARC1001873	F-0VARC1001873	4056	R-OVARC1001873	9048
	OVARC1001879	F-0VARC1001879	4057	R-0VARC1001879	9049
	OVARC1001880	F-0VARC1001880	4058	R-OVARC1001880	9050
50	OVARC1001883	F-0VARC1001883	4059	R-0VARC1001883	9051
	OVARC1001900	F-0VARC1001900	4060	R-OVARC1001900	9052
	OVARC1001901	F-0VARC1001901	4061	R-OVARC1001901	9053
	OVARC1001911	F-0VARC1001911	4062	R-OVARC1001911	9054
55	OVARC1001916	F-0VARC1001916	4063	R-0VARC1001916	9055

	OVARC1001928	F-0VARC1001928	4064	R-0VARC1001928	9056
	OVARC1001942	F-0VARC1001942	4065	R-0VARC1001942	9057
	OVARC1001943	F-0VARC1001943	4066	R-0VARC1001943	9058
5	OVARC1001949	F-0VARC1001949	4067	R-0VARC1001949	9059
	OVARC1001950	F-0VARC1001950	4068	R-0VARC1001950	9060
	OVARC1001987	F-0VARC1001987	4069	R-0VARC1001987	9061
	OVARC1001989	F-0VARC1001989	4070	R-0VARC1001989	9062
10	OVARC1002044	F-0VARC1002044	4071	R-0VARC1002044	9063
	OVARC1002050	F-0VARC1002050	4072	R-0VARC1002050	9064
	OVARC1002066	F-0VARC1002066	4073	R-0VARC1002066	9065
	OVARC1002082	F-0VARC1002082	4074	R-0VARC1002082	9066
15	OVARC1002107	F-0VARC1002107	4075	R-0VARC1002107	9067
75	OVARC1002112	F-0VARC1002112	4076		
	OVARC1002127	F-OVARC1002127,	4077	R-0VARC1002127	9068
	OVARC1002138	F-0VARC1002138	4078	R-0VARC1002138	9069
	OVARC1002143	F-0VARC1002143	4079	R-OVARC1002143	9070
20	OVARC1002156	F-0VARC1002156	4080	R-OVARC1002156	9071
	OVARC1002158	F-0VARC1002158	4081	R-0VARC1002158	9072
	0VARC1002165	F-0VARC1002165	4082	R-0VARC1002165	9073
	OVARC1002182	F-0VARC1002182	4083	R-0VARC1002182	9074
25	PLACE1000004	F-PLACE1000004	4084	R-PLACE1000004	9075
	PLACE1000005	F-PLACE1000005	4085	R-PLACE1000005	9076
	PLACE1000007	F-PLACE1000007	4086	R-PLACE1000007	9077
	PLACE1000014	F-PLACE1000014	4087	R-PLACE1000014	9078
30	PLACE1000031	F-PLACE1000031	4088	R-PLACE1000031	9079
	PLACE1000040	F-PLACE1000040	4089	R-PLACE1000040	9080
	PLACE1000048	F-PLACE1000048	4090	R-PLACE1000048	9081
	PLACE1000050	F-PLACE1000050	4091	R-PLACE1000050	9082
25	PLACE1000061	F-PLACE1000061	4092	R-PLACE1000061	9083
35	PLACE1000066	F-PLACE1000066	4093	R-PLACE1000066	9084
	PLACE1000078	F-PLACE1000078	4094	R-PLACE1000078	9085
	PLACE1000081	F-PLACE1000081	4095	R-PLACE1000081	9086
	PLACE1000094	F-PLACE1000094	4096	R-PLACE1000094	9087
40	PLACE1000133	F-PLACE1000133	4097	R-PLACE1000133	9088
	PLACE1000142	F-PLACE1000142	4098	R-PLACE1000142	9089
	PLACE1000184	F-PLACE1000184	4099	R-PLACE1000184	9090
	PLACE1000185	F-PLACE1000185	4100	R-PLACE1000185	9091
45	PLACE1000213	F-PLACE1000213	4101	R-PLACE1000213	9092
	PLACE1000214	F-PLACE1000214	4102	R-PLACE1000214	9093
	PLACE1000236	F-PLACE1000236	4103	R-PLACE1000236	9094
	PLACE1000246	F-PLACE1000246	4104	R-PLACE1000246	9095
50	PLACE1000292	F-PLACE1000292	4105	R-PLACE1000292	9096
	PLACE1000308	F-PLACE1000308	4106		
	PLACE1000332	F-PLACE1000332	4107	R-PLACE1000332	9097
	PLACE1000347	F-PLACE1000347	4108	R-PLACE1000347	9098
	PLACE1000374	F-PLACE1000374	4109	R-PLACE1000374	9099
55					

	PLACE1000380	F-PLACE1000380	4110	R-PLACE1000380	9100 -
	PLACE1000383	F-PLACE1000383	4111	R-PLACE1000383	9101
	PLACE1000401	F-PLACE1000401	4112	R-PLACE1000401	9102
5	PLACE1000406	F-PLACE1000406	4113	R-PLACE1000406	9103
	PLACE1000420	F-PLACE1000420	4114	R-PLACE1000420	9104
	PLACE1000421	F-PLACE1000421	4115	R-PLACE1000421	9105
	PLACE1000424	F-PLACE1000424	4116	R-PLACE1000424	9106
10	PLACE1000435	F-PLACE1000435	4117	R-PLACE1000435	9107
10	PLACE1000444	F-PLACE1000444	4118	R-PLACE1000444	9108
	PLACE1000453	F-PLACE1000453	4119	R-PLACE1000453	9109
	PLACE1000481	F-PLACE1000481	4120	R-PLACE1000481	9110
	PLACE1000491	F-PLACE1000492	4121	R-PLACE1000481	9111
15	PLACE1000492	F-PLACE1000540	4121		
				R-PLACE1000540	9112
	PLACE1000547	F-PLACE1000547,	4123	R-PLACE1000547	9113
	PLACE1000562	F-PLACE1000562	4124	R-PLACE1000562	9114
20	PLACE1000564	F-PLACE1000564	4125	R-PLACE1000564	9115
	PLACE1000583	F-PLACE1000583	4126	R-PLACE1000583	9116
	PLACE1000588	F-PLACE1000588	4127	R-nnnnnnnnnnn	9117
	PLACE1000596	F-PLACE1000596	4128	R-PLACE1000596	9118
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	PLACE4000465	F-PLACE4000465	5229	R-PLACE4000465	10169
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	PLACE4000548	F-PLACE4000548	5235	R-PLACE4000548	10173
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	Y79AA1000131	F-Y79AA1000131	5416	R-Y79AA1000131	10337
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	Y79AA1000420	F-Y79AA1000420	5433	R-Y79AA1000420	10354
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45	Y79AA1000480	F-Y79AA1000480	5435	R-Y79AA1000480	10356
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35	Y79AA1001145	F-Y79AA1001145	5472	R-Y79AA1001145	10390
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	Y79AA1001391	F-Y79AA1001391	5486	R-Y79AA1001391	10404
	Y79AA1001394	F-Y79AA1001394	5487	R-Y79AA1001394	10405
55	Y79AA1001402	F-Y79AA1001402	5488	R-Y79AA1001402	10406
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	Y79AA1001555	F-Y79AA1001555	5494	R-Y79AA1001555	10412
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	Y79AA1001613	F-Y79AA1001613	5499	R-Y79AA1001613	10416
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	Y79AA1001781	F-Y79AA1001781	5507	R-Y79AA1001781	10424
	Y79AA1001805	F-Y79AA1001805	5508	R-nnnnnnnnnnn	10425
25	Y79AA1001827	F-Y79AA1001827	5509	R-Y79AA1001827	10426
	Y79AA1001846	F-Y79AA1001846	5510	R-Y79AA1001846	10427
	Y79AA1001848	F-Y79AA1001848	5511	R-Y79AA1001848	10428
	Y79AA1001866	F-Y79AA1001866	5512	R-Y79AA1001866	10429
30	Y79AA1001874	F-Y79AA1001874	5513	R-Y79AA1001874	10430
	Y79AA1001875	F-Y79AA1001875	5514	R-Y79AA1001875	10431
	Y79AA1001923	F-Y79AA1001923	5515	R-Y79AA1001923	10432
	Y79AA1001963	F-Y79AA1001963	5516		
35	Y79AA1002027	F-Y79AA1002027	5517	R-Y79AA1002027	10433
	Y79AA1002083	F-Y79AA1002083	5518	R-Y79AA1002083	10434
	Y79AA1002089	F-Y79AA1002089	5519	R-Y79AA1002089	10435
	Y79AA1002093	F-Y79AA1002093	5520	R-Y79AA1002093	10436
40	Y79AA1002103	F-Y79AA1002103	5521	R-Y79AA1002103	10437
40	Y79AA1002115	F-Y79AA1002115	5522	R-Y79AA1002115	10438
	Y79AA1002125	F-Y79AA1002125	5523	R-Y79AA1002125	10439
	Y79AA1002139	F-Y79AA1002139	5524	R-Y79AA1002139	10440
	Y79AA1002204	F-Y79AA1002204	5525	R-Y79AA1002204	10441
45	Y79AA1002208	F-Y79AA1002208	5526	R-nnnnnnnnnnn	10442
	Y79AA1002209	F-Y79AA1002209	5527	R-Y79AA1002209	10443
	Y79AA1002210	F-Y79AA1002210	5528	R-Y79AA1002210	10444
	Y79AA1002211	F-Y79AA1002211	5529	R-Y79AA1002211	10445
50	Y79AA1002220	F-Y79AA1002220	5530	R-Y79AA1002220	10446
	Y79AA1002229	F-Y79AA1002229	5531	R-Y79AA1002229	10447
	Y79AA1002234	F-Y79AA1002234	5532	R-Y79AA1002234	10448
	Y79AA1002246	F-Y79AA1002246	5533	R-Y79AA1002246	10449
55	Y79AA1002258	F-Y79AA1002258	5534	R-Y79AA1002258	10450
	Y79AA1002298	F-Y79AA1002298	5535	R-Y79AA1002298	10451

	Y79AA1002307	F-Y79AA1002307	5536	R-Y79AA1002307	10452
	Y79AA1002311	F-Y79AA1002311	5537	R-Y79AA1002311	10453
5	Y79AA1002351	F-Y79AA1002351	5538	R-Y79AA1002351	10454
· ·	Y79AA1002361	F-Y79AA1002361	5539	R-Y79AA1002361	10455
	Y79AA1002399	F-Y79AA1002399	5540	R-Y79AA1002399	10456
	Y79AA1002407	F-Y79AA1002407	5541	R-Y79AA1002407	10457
10	Y79AA1002416	F-Y79AA1002416	5542	R-Y79AA1002416	10458
	Y79AA1002431	F-Y79AA1002431	5543	R-Y79AA1002431	10459
	Y79AA1002433	F-Y79AA1002433	5544	R-nnnnnnnnnnn	10460
	Y79AA1002472	F-Y79AA1002472	5545	R-Y79AA1002472	10461
15	Y79AA1002482	F-Y79AA1002482	5546	R-Y79AA1002482	10462
	Y79AA1002487	F-Y79AA1002487	5547	R-Y79AA1002487	10463

Table 2

	name of	name of	SEQ ID	name of	SEQ ID
	clone	5'-end	of 5'-end	3'-end	of 3'-end
		sequence	sequence	sequence	sequence
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	HEMBA1000497	F-HEMBA100049	7 16111	R-HEMBA100049	97 16165
	HEMBA1001750	F-HEMBA100175	0 16112	R-HEMBA10017	50 16166
15	HEMBA1003854	F-HEMBA100385	4 16113	R-HEMBA10038	54 16167
70	HEMBA1004193	F-HEMBA100419	3 16114	R-HEMBA100419	93 16168
	HEMBA1004860	F-HEMBA100486	0 16115	R-HEMBA100486	60 16169
	HEMBA1005572	F-HEMBA100557	2 16116	R-HEMBA10055	72 16170
20	HEMBA1006038	F-HEMBA100603	8 16117	R-HEMBA10060:	38 16171
	HEMBA1006092	F-HEMBA100609	2 16118	R-HEMBA100609	92 16172
	HEMBA1006406	F-HEMBA100640	6 16119	R-HEMBA10064	06 16173
	HEMBA1006650	F-HEMBA100665	0 16120	R-HEMBA10066	50 16174
25	HEMBA1006812	F-HEMBA100681	2 16121	R-HEMBA10068	12 16175
	HEMBB1000672	F-HEMBB100067	2 16122	R-HEMBB10006	72 16176
	HEMBB1001197	F-HEMBB100119	7 16123	R-HEMBB100119	97 16177
	HEMBB1001871	F-HEMBB100187	1 16124	R-HEMBB10018	71 16178
30	MAMMA1001252	F-MAMMA100125	2 16125	R-MAMMA10012	52 16179
	MAMMA1002094	F-MAMMA100209	4 16126	R-MAMMA10020	94 16180
	NT2RM4000634	F-NT2RM400063	4 16127	R-NT2RM40006	34 16181
	NT2RM4000657	F-NT2RM400065	7 16128	R-NT2RM40006	57 16182
35	NT2RM4000783	F-NT2RM400078	3 16129	R-NT2RM40007	83 16183
	NT2RM4000857	F-NT2RM400085	7 16130	R-NT2RM40008	57 16184
	NT2RM4001178	F-NT2RM400117	8 16131	R-NT2RM40011	78 16185
40	NT2RM4002420	F-NT2RM400242	0 16132	R-NT2RM40024	20 16186
	NT2RP2000198	F-NT2RP200019	8 16133	R-NT2RP20001	98 16187
	NT2RP2000551	F-NT2RP200055		R-NT2RP20005	
	NT2RP2000660	F-NT2RP200066	0 16135	R-NT2RP20006	60 16189

	NT2RP2001214	F-NT2RP2001214	16136	R-NT2RP2001214	16190
	NT2RP2001460	F-NT2RP2001460	16137	R-NT2RP2001460	16191
5	NT2RP2001756	F-NT2RP2001756	16138	R-NT2RP2001756	16192
•	NT2RP2002056	F-NT2RP2002056	16139	R-NT2RP2002056	16193
	NT2RP2002677	F-NT2RP2002677	16140	R-NT2RP2002677	16194
	NT2RP2002755	F-NT2RP2002755	16141	R-NT2RP2002755	16195
10	NT2RP2002843	F-NT2RP2002843	16142	R-NT2RP2002843	16196
	NT2RP2003101	F-NT2RP2003101	16143	R-NT2RP2003101	16197
	NT2RP2003799	F-NT2RP2003799	16144	R-NT2RP2003799	16198
	NT2RP2004095	F-NT2RP2004095	16145	R-NT2RP2004095	16199
15	NT2RP2004732	F-NT2RP2004732	16146	R-NT2RP2004732	16200
	NT2RP2004920	F-NT2RP2004920	16147	R-NT2RP2004920	16201
	NT2RP2005454	F-NT2RP2005454	16148	R-NT2RP2005454	16202
20	NT2RP2005776	F-NT2RP2005776	16149	R-NT2RP2005776	16203
20	NT2RP2005806	F-NT2RP2005806	16150	R-NT2RP2005806	16204
	NT2RP2005882	F-NT2RP2005882	16151	R-NT2RP2005882	16205
	NT2RP3001282	F-NT2RP3001282	16152	R-NT2RP3001282	16206
25	NT2RP3001723	F-NT2RP3001723	16153	R-NT2RP3001723	16207
	NT2RP3002099	F-NT2RP3002099	16154	R-NT2RP3002099	16208
	NT2RP3003155	F-NT2RP3003155	16155	R-NT2RP3003155	16209
	NT2RP3004028	F-NT2RP3004028	16156	R-NT2RP3004028	16210
30	0VARC1000008	F-0VARC1000008	16157	R-0VARC1000008	16211
	0VARC1000724	F-0VARC1000724	16158	R-0VARC1000724	16212
	0VARC1000751	F-0VARC1000751	16159	R-0VARC1000751	16213
	OVARC1001029	F-OVARC1001029	16160	R-0VARC1001029	16214
35	PLACE1000814	F-PLACE1000814	16161	R-PLACE1000814	16215
	PLACE1003030	F-PLACE1003030	16162	R-PLACE1003030	16216
	PLACE1005549	F-PLACE1005549	16163	R-PLACE1005549	16217
40	PLACE1007218	F-PLACE1007218	16164	R-PLACE1007218	16218

[0019] Furthermore, the present invention relates to the use of the above primers, as described below.

- (4) A polynucleotide which can be synthesized with the primer set of (2) or (3).
- (5) A polynucleotide comprising a coding region in the polynucleotide of (4).
- (6) A substantially pure protein encoded by polynucleotide of (4).
- (7) A partial peptide of the protein of (6).

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[0020] In addition, the present invention comprises a polynucleotide described below and a protein encoded by the polynucleotide.

- (8) An isolated polynucleotide selected from the group consisting of
 - (a) a polynucleotide comprising a coding region of the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
 - (b) a polynucleotide comprising a nucleotide sequence encoding a protein comprising the amino acid sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351;
 - (c) a polynucleotide comprising a nucleotide sequence encoding a protein comprising an amino acid sequence

selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, in which one or more amino acids are substituted, deleted, inserted, and/or added, wherein said protein is functionally equivalent to the protein comprising said amino acid sequence selected from the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351;

- (d) a polynucleotide that hybridizes with a polynucleotide comprising a nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351, and that comprises a nucleotide sequence encoding a protein functionally equivalent to the protein encoded by the nucleotide sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351;
- (e) a polynucleotide comprising a nucleotide sequence encoding a partial amino acid sequence of a protein encoded by the polynucleotide of (a) to (d);
- (f) a polynucleotide comprising a nucleotide sequence with at least 70% identity to the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351.
- (9). A substantially pure protein encoded by the polynucleotide of (8).
- (10) An antibody against the protein or peptide of any one of (6), (7), and (9).
- (11) A vector comprising the polynucleotide of (5) or (8).

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- (12) A transformant carrying the polynucleotide of (5) or (8), or the vector of (11).
- (13) A transformant expressively carrying the polynucleotide of (5) or (8), or the vector of (11).
- (14) A method for producing the protein or peptide of any one of (6), (7), and (9), comprising culturing the transformant of (13) and recovering the expression product.
- (15) An oligonucleotide comprising the nucleotide sequence set forth in any one of the SEQ ID NOs in Tables 350 and 351 or the nucleotide sequence complementary to the complementary strand thereof, wherein said oligonucleotide comprises 15 nucleotides or more.
- (16) Use of the oligonucleotide of (15) as a primer for synthesizing a polynucleotide.
- (17) Use of the oligonucleotide of (15) as a probe for detecting a gene.
- (18) An antisense polynucleotide against the polynucleotide of (8), or the portion thereof.
- (19) A method for synthesizing a polynucleotide, the method comprising:
 - a) synthesizing a complementary strand using a cDNA library as a template, and using the primer set of (2) or (3), or the primer of (16); and
 - b) recovering the synthesized product.
- (20) The method of (19), wherein the cDNA library is obtainable by oligo-capping method.
- (21) The method of (19), wherein the complementary strand is obtainable by PCR.
- (22) A method for detecting the polynucleotide of (8), the method comprising:
 - a) incubating a target polynucleotide with the oligonucleotide of (15) under the conditions where hybridization occurs, and
 - b) detecting the hybridization of the target polynucleotide with the oligonucleotide of (15).
- (23) A database of polynucleotides and/or proteins, the database comprising information on at least one sequence selected from the nucleotide sequences set forth in the SEQ ID NOs in Tables 350 and 351 and/or the amino acid sequences set forth in the SEQ ID NOs in Tables 350 and 351, or a medium on which the database is stored.
- 45 [0021] Any patents, patent applications, and publications cited herein are incorporated by reference.

BRIEF DESCRIPTION OF THE DRAWINGS

- [0022] Figure 1 shows the restriction maps of vectors pME18SFL3 and pUC19FL3.
- [0023] Figure 2 shows the reproducibility of gene expression analysis. The respective intensities of gene expression observed in independent set of experiments are plotted in the vertical axis as well as in the horizontal axis.
- [0024] Figure 3 shows the detection limit in gene expression analysis. The intensity of expression is shown in the vertical axis and the concentration (µg/ml) of probe used is shown in the horizontal axis.

DETAILED DESCRIPTION OF THE INVENTION

[0025] Herein, "polynucleotide" is defined as a molecule in which multiple nucleotides are polymerized. There are no limitations in the number of the polymerized nucleotides. In case that the polymer contains relatively low number

of nucleotides, it is also described as an "oligonucleotide". The polynucleotide or the oligonucleotide of the present invention can be a natural or chemically synthesized product. Alternatively, it can be synthesized using a template DNA by an enzymatic reaction such as PCR.

[0026] All the cDNA provided by the invention are full-length cDNA. Herein, a "full-length cDNA" is defined as a cDNA which contains both ATG codon (the translation start site) and the stop codon. Accordingly, the untranslated regions, which are originally found in the upstream or downstream of the protein coding region in natural mRNA, may or may not be contained

[0027] An "isolated polynucleotide" is a polynucleotide the structure of which is not identical to that of any naturally occurring nucleic acid or to that of any fragment of a naturally occurring genomic nucleic acid spanning more than three separate genes. The term therefore covers, for example,

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- (a) a DNA which has the sequence of part of a naturally occurring genomic DNA molecule but is not flanked by both of the coding sequences that flank that part of the molecule in the genome of the organism in which it naturally occurs;
- (b) a nucleic acid incorporated into a vector or into the genomic DNA of a prokaryote or eukaryote in a manner such that the resulting molecule is not identical to any naturally occurring vector or genomic DNA;
- (c) a separate molecule such as a cDNA, a genomic fragment, a fragment produced by polymerase chain reaction (PCR), or a restriction fragment; and
- (d) a recombinant nucleotide sequence that is part of a hybrid gene, i.e., a gene encoding a fusion protein. Specifically excluded from this definition are nucleic acids present in mixtures of different (i) DNA molecules, (ii) transfected cells, or (iii) cell clones: e.g., as these occur in a DNA library such as a cDNA or genomic DNA library.

[0028] The term "substantially pure" as used herein in reference to a given polypeptide means that the protein or polypeptide is substantially free from other biological macromolecules. The substantially pure protein or polypeptide is at least 75% (e.g., at least 80, 85, 95, or 99%) pure by dry weight. Purity can be measured by any appropriate standard method, for example, by column chromatography, polyacrylamide gel electrophoresis, or HPLC analysis.

[0029] All the clones (5602 clones) of the present invention are novel and encode the full-length proteins. All the clones were prepared by oligo capping method, which can achieve cDNA cloning with high fullness ratio. The cDNA clones were selected by using ATGprl score as an index of the fullness ratio at the 5'-end, based on the sequence features of the 5'-end sequences. Selection was further carried out by searching GenBank database for EST sequences homologous to 5'-end sequence of each clone by BLAST [S.F. Altschul, W. Gish, W. Miller, E.W. Myers & D.L. Lipman J. Mol. Biol., 215:403-410 (1990); W. Gish, & D.J. States, Nature Genet., 3:266-272 (1993)] and by considering the number of matching (identical) EST sequences or the number of continuous amino acids in the 5'-end sequence initiated from the initiation codon.

[0030] Moreover, the clones were turn out to be not identical to any of the known human mRNA (namely novel) by homology search using the 5'-end sequence.

[0031] The primers of the present invention, which are used for synthesizing full-length cDNA, are selected from the group comprising SEQ ID NO: 1-5547 (5'-primer), or SEQ ID NO: 5548-10463 (3'-primer). Further, the primers of the present invention, which are used for synthesizing full-length cDNA, are selected from SEQ ID NO: 16111-16164 (5'-primer), or SEQ ID NO: 16165-16218 (3'-primer). Some of the nucleotides include a known EST as its part. However, the primers of the present invention are novel in terms that the primers enable to synthesize full-length cDNA. Because the known ESTs lack important information on what part of cDNA the ESTs correspond to, it is impossible to design primers on the basis of the ESTs.

[0032] All the full-length cDNA of the present invention can be synthesized using a primer set comprising the nucleotide sequences selected from both the 5'-and 3'-end sequences, or a set comprising a primer based on the 5'-end sequence and an oligo-dT primer, by a method such as PCR (Current protocols in Molecular Biology (1987) Ausubel et al. edit, John Wiley & Sons, Section 6.1-6.4).

[0033] Specifically, PCR can be performed using an oligonucleotide that has 15 nucleotides longer, and specifically hybridizes with the complementary strand of the polynucleotide that contains the nucleotide sequence selected from the 5'-end sequences shown in Table 1 and 2 (SEQ ID NO: 1-5547, or SEQ ID NO: 16111-16164), and an oligo-dT primer as a 5'-, and 3'-primer, respectively. The length of the primers is usually 15-100 bp, and favorably between 15-35 bp. In case of LA PCR, which is described below, the primer length of 25-35 bp may provide a good result.

[0034] A method to design a primer that enables a specific amplification based on the given nucleotide sequence is known to those skilled in the art (Current Protocols in Molecular Biology, Ausubel et al. edit, (1987) John Wiley & Sons, Section 6.1-6.4). In designing a primer based on the 5'-end sequence, the primer is designed so as that, in principle, the amplification products will include the translation start site. Accordingly, in case that a given 5'-end nucleotide sequence is the 5'- untranslated region (5'UTR), any part of the sequence can be used as a 5'-primer as far as the specificity toward the target cDNA is insured. The translation start site can be predicted using a known method such

as the ATGpr as described below.

[0035] When synthesizing a polynucleotide, the target nucleotide sequence to be amplified can extend to several thousand bp in some cDNA. However, it is possible to amplify such a long nucleotides by using such as LA PCR (Long and Accurate PCR). It is advantageous to use LA PCR when synthesizing long DNA. In LA PCR, in which a special DNA polymerase having 3' \rightarrow 5' exonuclease activity is used, misincorporated nucleotides can be removed. Accordingly, accurate synthesis of the complementary strand can be achieved even with a long nucleotide sequence. By using LA PCR, it is reported that amplification of a nucleotide with 20 kb longer can be achieved under desirable condition (Takeshi Hayashi (1996) Jikken-Igaku Bessatsu, "Advanced Technologies in PCR" Youdo-sha).

[0036] A template DNA for synthesizing the cDNA of the present invention can be obtained by using cDNA libraries that are prepared by various methods. The full-length cDNA clones obtained here are those with high fullness ratio, which were obtained using a combination of (1) a method to prepare a full-length-enriched cDNA library using the oligo-capping method, and (2) an estimation system for fullness using the 5'-end sequence (selection based on the estimation by the ATGpr after removing clones that are not-full-length compared to the ESTs). However, it is possible to easily obtain a full-length cDNA by using the primers that are provided by the present invention, not by the above described specialized method.

The problem with the cDNA libraries prepared by the known methods or commercially available is that mRNA contained in the libraries has very low fullness ratio. Thus, it is difficult to screen full-length cDNA clone directly from the library using ordinary cloning methods. The present invention has revealed a primer that is capable of synthesizing a full-length cDNA. If provided with primers, it is possible to synthesize a target full-length cDNA by using enzymatic reactions such as PCR. In particular, a full-length-enriched cDNA library, synthesized by methods such as oligo-capping, is desirable to synthesize a full-length cDNA with more reliability.

[0037] The 5'-end sequence of the full-length cDNA clones of the invention can be used to isolate the regulatory element of transcription including the promoter on the genome. By the spring of the year 2000, a rough draft of the human genome (analysis of human genomic sequence with lower accuracy), which covers 90% of the genome, is planned to be accomplished, and by the year 2003, analysis of the entire human genomic sequence is going to be finished. However, it is hard to analyze with software the transcription start sites on the human genome, in which long introns exist. By contrast, it is easy to specify the transcription start site on the genomic sequence using the 5'-end sequence of the full-length cDNA clone, thus it is easy to obtain the genomic region involved in transcription regulation, which includes the promoter that is contained in the upstream of the transcription start site.

[0038] The full-length cDNAs cloned in the present invention are classified into 13 groups, based on the data such as ATGpr1 score, by which the fullness ratio can be evaluated. Specifically, the 13 groups consist of; the below-mentioned groups (1)-(3), containing 3690 clones (Table 9), and the group (12), containing 3 clones, wherein ATGpr1 (score defined in the ATGpr program) is higher than 0.3; and the below-mentioned groups (4)-(11), containing 1857 clones (Table 10), and the group (13), containing 52 clones, wherein, although ATGpr1 is 0.3 or less, the clones are judged to be full-length from various viewpoints. Names of the clones belonging to the groups (1)-(13) are as indicated in Examples or below.

(1) 1516 clones

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Among the 3690 clones that have the maximal ATGpr1 score higher than 0.3, 1516 clones are novel full-length clones, in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

(2)377 clones

Among the 3690, 377 clones are novel full-length clones, in which the number of human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

(3) 1797 clones

Among the 3690, 1797 clones are novel full-length clones, in which the number of human EST having identical sequence at the 5'-end is not more than 20 (except the clones described above).

(4) 453 clones

Among the 1857 clones in which the maximal ATGpr1 score is 0.3 or less, the following 453 clones are estimated to be novel full-length clones since the clones have the maximal score 0.3 or more in the ATGpr2, and at least either of the sequences of their 5'- and 3'-ends, or both are not identical to those of any human EST. The ATGpr2 score is determined by using the ATGpr program with neglecting the information of the frequency of the six nucleotides contained within the sequence between the ATG codon and the stop codon (the maximal length is 300 nucleotides from the ATG codon) (Salamo A.A., Nishikawa T., and Swindells M.B. (1998) Bioinformatics, 14: 384-390; http://www.hri.co.jp/atgpr/). The ATGpr program for calculating the ATGpr2 score is described as the

ATGpr2 program in the followings.

(5) 24 clones

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Among the 1857 clones, 24 clones are estimated to be full-length since their maximal ATGpr2 scores are higher than 0.3, and also novel, though they have low scores in ATGpr1 program, in which the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

(6) 65 clones

Among the 1857 clones, 65 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

(7) 32 clones

Among the 1857 clones, 32 clones are estimated to be full-length since, though they have low scores in both programs, ATGpr1 and ATGpr2, the scores are the maximum in comparison to those of the other clones in the same cluster (at least two clones). The clones are also novel, if the number of the human EST having identical sequence at both 5'- and 3'-ends is 1 to 5.

(8) 36 clones

Among the 1857 clones, 36 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST.

(9) 81 clones

Among the 1857 clones, 81 clones are full-length, which were selected by assembling the sequences of the other clones or human EST, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if the number of the human EST having identical sequence at the 5'-end is not more than 20 (other than the clones in which at least either of the sequences of the 5'- and 3'-ends, or both are not identical to those of any human EST).

(10) 938 clones

Among the 1857 clones, 938 clones are estimated to be full-length according to the fullness ratio shown in Table 4, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 5'-end is not identical to those of any human EST.

(11) 228 clones

Among the 1857 clones, 228 clones are estimated to be full-length according to the fullness ratio shown in Table 7, although they have low scores in both programs, ATGpr1 and ATGpr2. The clones are also novel, if at least the sequence of the 3'-end is not identical to those of any human EST.

(12) 3 clones

Three clones, HEMBA1006812, HEMBB1001871, and NT2RP3001282, whose maximal ATGpr1 values are higher than 0.3, are full-length and novel clones whose 5'-end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 100 amino acids or more.

(13) 52 clones

The following 52 clones, which have maximal ATGpr1 values of 0.3 or less, are full-length with the fullness ratios shown in Table 4 although the fullness ratios are low:

HEMBA1000497,	HEMBA1001750,	HEMBA1003854,	HEMBA1004193,	HEMBA1004860,	HEMBA1005572,
HEMBA1006038,	HEMBA1006092,	HEMBA1006406,	HEMBA1006650,	HEMBB1000672,	HEMBB1001197,
MAMMA1001252,	MAMMA1002094,	NT2RM4000634,	NT2RM4000657,	NT2RM4000783,	NT2RM4000857,
NT2RM4001178,	NT2RM4002420,	NT2RP2000198,	NT2RP2000551,	NT2RP2000660,	NT2RP2001214,
NT2RP2001460,	NT2RP2001756,	NT2RP2002056,	NT2RP2002677,	NT2RP2002755,	NT2RP2002843,
NT2RP2003101,	NT2RP2003799,	NT2RP2004095,	NT2RP2004732,	NT2RP2004920,	NT2RP2005454,
NT2RP2005776,	NT2RP2005806,	NT2RP2005882,	NT2RP3001723,	NT2RP3002099,	NT2RP3003155.
NT2RP3004028,	OVARC1000008,	OVARC1000724,	OVARC1000751,	OVARC1001029.	PLACE1000814.

PLACE1003030, PLACE1005549, PLACE1007218, NT2RP4002298.

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Moreover, the clones are novel clones whose 5' -end sequences presumably contain a coding region which is initiated with ATG codon and which encodes 50 amino acids or more. Among them, the following 20 clones is predicted to contain a coding region with 100 amino acids or more and should encode proteins:

HEMBA1000497, HEMBA1003854, HEMBA1004193, NT2RM4000657, NT2RM4001178, NT2RP2001756, NT2RP2002677, NT2RP2002755, NT2RP2002843, NT2RP2004095, NT2RP2004920, NT2RP2005806, NT2RP3002099, NT2RP3003155, OVARC1000724, OVARC1001029, PLACE1000814, PLACE1003030, PLACE1005549, PLACE1007218.

[0039] The protein encoded by the polynucleotide of the invention can be prepared as a recombinant protein or as a natural protein. For example, the recombinant protein can be prepared by inserting the polynucleotide encoding the protein of the invention into a vector, introducing the vector into an appropriate host cell and purifying the protein expressed within the transformed host cell, as described below. In contrast, the natural protein can be prepared, for example, by utilizing an affinity column to which an antibody against the protein of the invention (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 16.1-16.19) is attached. The antibody used for affinity purification may be either a polyclonal antibody, or a monoclonal antibody. Alternatively, in vitro translation (See, for example, "On the fidelity of mRNA translation in the nuclease-treated rabbit reticulocyte lysate system." Dasso M.C., and Jackson R.J. (1989) Nucleic Acids Res. 17: 3129-3144) may be used for preparing the protein of the invention. [0040] Proteins functionally equivalent to the proteins of the present invention can be prepared based on the activities, which were clarified in the above-mentioned manner, of the proteins of the present invention. Using the biological activity possessed by the protein of the invention as an index, it is possible to verify whether or not a particular protein is functionally equivalent to the protein of the invention by examining whether or not the protein has said activity.

[0041] Proteins functionally equivalent to the proteins of the present invention can be prepared by those skilled in the art, for example, by using a method for introducing mutations into an amino acid sequence of a protein (for example, site-directed mutagenesis (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 8.1-8.5). Besides, such proteins can be generated by spontaneous mutations. The present invention comprises the proteins having one or more amino acids substitutions, deletions, insertions and/or additions in the amino acid sequences of the proteins of the present invention (Tables 350 and 351), as far as the proteins have the equivalent functions to those of the proteins identified in the present Examples described later.

[0042] There are no limitations in the number and sites of amino acid mutations, as far as the proteins maintain the functions thereof. The number of mutations is typically 30% or less, or 20% or less, or 10% or less, preferably within 5% or less, or 3% or less of the total amino acids, more preferably within 2% or less or 1 % or less of the total amino acids. From the viewpoint of maintaining the protein function, it is preferable that a substituted amino has a similar property to that of the original amino acid. For example, Ala, Val, Leu, Ile, Pro, Met, Phe and Trp are assumed to have similar properties to one another because they are all classified into a group of non-polar amino acids. Similarly, substitution can be performed among non-charged amino acid such as Gly, Ser, Thr, Cys, Tyr, Asn, and Gln, acidic amino acids such as Asp and Glu, and basic amino acids such as Lys, Arg, and His.

[0043] In addition, proteins functionally equivalent to the proteins of the present invention can be isolated by using techniques of hybridization or gene amplification known to those skilled in the art. Specifically, using the hybridization technique (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)), those skilled in the art can usually isolate a DNA highly homologous to the DNA encoding the protein identified in the present Example based on the identified nucleotide sequence (Tables 350 and 351) or a portion thereof and obtain the functionally equivalent protein from the isolated DNA. The present invention include proteins encoded by the DNAs hybridizing with the DNAs encoding the proteins identified in the present Example, as far as the proteins are functionally equivalent to the proteins identified in the present Example. Organisms from which the functionally equivalent proteins are isolated are illustrated by vertebrates such as human, mouse, rat, rabbit, pig and bovine, but are not limited to these animals.

[0044] Washing conditions of hybridization for the isolation of DNAs encoding the functionally equivalent proteins are usually " $1 \times SSC$, 0.1% SDS, $37^{\circ}C$ "; more stringent conditions are " $0.5 \times SSC$, 0.1% SDS, $42^{\circ}C$ "; and still more stringent conditions are " $0.1 \times SSC$, 0.1% SDS, $65^{\circ}C$ ". Alternatively, the following conditions can be given as hybridization conditions of the present invention. Namely, conditions in which the hybridization is done at " $6 \times SSC$, 40% Formamide, $25^{\circ}C$ ", and the washing at " $1 \times SSC$, $55^{\circ}C$ " can be given. More preferable conditions are those in which the hybridization is done at " $6 \times SSC$, 40% Formamide, $37^{\circ}C$ ", and the washing at " $0.2 \times SSC$, $55^{\circ}C$ ". Even more preferable are those in which the hybridization is done at " $6 \times SSC$, 50% Formamide, $37^{\circ}C$ ", and the washing at " $0.1 \times SSC$, $62^{\circ}C$ ". The more stringent the conditions of hybridization are, the more frequently the DNAs highly homologous to the probe sequence are isolated. Therefore, it is preferable to conduct hybridization under stringent conditions. Examples of stringent conditions in the present invention are, washing conditions of " $0.5 \times SSC$, 0.1% SDS, $42^{\circ}C$ ", or alternatively, hybridization conditions of " $6 \times SSC$, 40% Formamide, $37^{\circ}C$ ", and the washing at " $0.2 \times SSC$, $55^{\circ}C$ ". However, the above-mentioned combinations of SSC, SDS and temperature conditions are indicated just as examples.

Those skilled in the art can select the hybridization conditions with similar stringency to those mentioned above by properly combining the above-mentioned or other factors (for example, probe concentration, probe length and duration of hybridization reaction) that determines the stringency of hybridization.

[0045] The amino acid sequences of proteins isolated by using the hybridization techniques usually exhibit high homology to those of the proteins of the present invention, which are shown in Tables 350 and 351. The present invention encompasses a polynucleotide comprising a nucleotide sequence that has a high identity to the nucleotide sequence of claim 8 (a).

Furthermore, the present invention encompasses a peptide, or protein comprising an amino acid sequence that has a high identity to the amino acid sequence encoded by the polynucleotide of claim 8 (b). The term "high identity" indicates sequence identity of at least 40% or more;

preferably 60% or more; and more preferably 70% or more. Alternatively, more preferable is identity of 90% or more, or 93% or more, or 95% or more, furthermore, 97% or more, or 99% or more. The identity can be determined by using the BLAST search algorithm.

[0046] With the gene amplification technique (PCR) (Current Protocols in Molecular Biology, edit, Ausubel et al., (1987) John Wiley & Sons, Section 6.3-6.4)) using primers designed based on the nucleotide sequence (Tables 350 and 351) or a portion thereof identified in the present Example, it is possible to isolate a DNA fragment highly homologous to the polynucleotide sequence or a portion thereof and to obtain functionally equivalent protein to a particular protein identified in the present Example based on the isolated DNA fragment.

[0047] The "percent identity" of two amino acid sequences or of two nucleic acids is determined using the algorithm of Karlin and Altschul (Proc. Natl. Acad. Sei. USA 87:2264-2268, 1990), modified as in Karlin and Altschul (Proc. Natl. Acad. Sei. USA 90:5873-5877, 1993). Such an algorithm is incorporated into the BLASTN and BLASTX programs of Altschul et al. (J. Mol. Biol.215:403-410, 1990). BLAST nucleotide searches are performed with the BLASTN program, score = 100, wordlength = 12. BLAST protein searches are performed with the BLASTX program, score = 50, wordlength = 3. When gaps exist between two sequences, Gapped BLAST is utilized as described in Altschul et al. (Nucleic Acids Res.25:3389-3402,1997). When utilizing BLAST and Gapped BLAST programs, the default parameters of the respective programs (e.g., BLASTX and BLASTN) are used. See http://www.ncbi.nlm.nih.gov.

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[0048] The present invention also includes a partial peptide of the proteins of the invention. The partial peptide comprises a protein generated as a result that a signal peptide has been removed from a secretory protein. If the protein of the present invention has an activity as a receptor or a ligand, the partial peptide may function as a competitive inhibitor of the protein and may bind to the receptor (or ligand). In addition, the present invention comprises an antigen peptide for raising antibodies. For the peptides to be specific for the protein of the invention, the peptides comprise at least 7 amino acids, preferably 8 amino acids or more, more preferably 9 amino acids or more, and even more preferably 10 amino acids or more. The peptide can be used for preparing antibodies against the protein of the invention, or competitive inhibitors of them, and also screening for a receptor that binds to the protein of the invention. The partial peptides of the invention can be produced, for example, by genetic engineering methods, known methods for synthesizing peptides, or digesting the protein of the invention with an appropriate peptidase.

[0049] The present invention also relates to a vector into which the DNA of the invention is inserted. The vector of the invention is not limited as long as it contains the inserted DNA stably. For example, if *E. coli* is used as a host, vectors such as pBluescript vector (Stratagene) are preferable as a cloning vector. To produce the protein of the invention, expression vectors are especially useful. Any expression vector can be used as far as it is capable of expressing the protein *in vitro*, in *E. coli*, in cultured cells, or *in vivo*. For example, pBEST vector (Promega) is preferable for *in vitro* expression, pET vector (Invitrogen) for *E. coli*, pME18S-FL3 vector (GenBank Accession No. AB009864) for cultured cells, and pME18S vector (Mol. Cell. Biol. (1988) 8: 466-472) for *in vivo* expression. To insert the DNA of the invention, ligation utilizing restriction sites can be performed according to the standard method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

[0050] The present invention also relates to a transformant carrying the vector of the invention. Any cell can be used as a host into which the vector of the invention is inserted, and various kinds of host cells can be used depending on the purposes. For strong expression of the protein in eukaryotic cells, COS cells or CHO cells can be used, for example. Introduction of the vector into host cells can be performed, for example, by calcium phosphate precipitation method, electroporation method (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 9.1-9.9), lipofectamine method (GIBCO-BRL), or microinjection method, etc.

[0052] The primer of the present invention can be used for synthesizing full-length cDNA, and also for the detection and/or diagnosis of the abnormality of the protein of the invention encoded by the full-length cDNA. For example, by utilizing polymerase chain reaction (genomic DNA-PCR, or RT-PCR) using the primer of the invention, DNA encoding the protein of the invention can be amplified. It is also possible to obtain the regulatory region of expression in the 5'-upstream by using PCR or hybridization since the transcription start site within the genomic sequence can be easily specified based on the 5'-end sequence of the full-length cDNA. The obtained genomic region can be used for detection and/or diagnosis of the abnormality of the sequence by RFLP analysis, SSCP, or direct sequencing.

[0053] Furthermore, the "polynucleotide having a length of at least 15 nucleotides, comprising a nucleotide sequence that is complementary to a polynucleotide comprising the nucleotide sequence set forth in any one of SEQ ID NOs in Tables 350 and 351, or its complementary strand" includes an antisense polynucleotide for suppressing the expression of the protein of the invention. To exert the antisense effect, the antisense polynucleotide has a length of at least 15 bp or more, for example, 50 bp or more, preferably 100 bp or more, and more preferably 500 bp or more, and has a length of usually 3000 bp or less and preferably 2000 bp or less. The antisense DNA can be used in the gene therapy of the diseases that are caused by the abnormality of the protein of the invention (abnormal function or abnormal expression). Said antisense DNA can be prepared, for example, by the phosphorothioate method ("Physicochemical properties of phosphorothioate oligodeoxynucleotides." Stein (1988) Nucleic Acids Res. 16: 3209-3221) based on the nucleotide sequence of the DNA encoding the protein (for example, the DNA set forth in any one of SEQ ID NOs in Tables 350 and 351).

[0054] The polynucleotide or antisense DNA of the present invention can be used in gene therapy, for example, by administrating it into a patient by the in vivo or ex vivo method with virus vectors such as retrovirus vectors, adenovirus vectors, and adeno-associated virus vectors, or non-virus vectors such as liposome.

[0055] The present invention also relates to antibodies that bind to the protein of the invention.

There are no limitations in the form of the antibodies of the invention. They include polyclonal antibodies, monoclonal antibodies, or their portions that can bind to the protein of the invention. They also include antibodies of all classes. Furthermore, special antibodies such as humanized antibodies are also included.

[0056] The polyclonal antibody of the invention can be obtained according to the standard method by synthesizing an oligopeptide corresponding to the amino acid sequence and immunizing rabbits with the peptides (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.12-11.13). The monoclonal antibody of the invention can be obtained according to the standard method by purifying the protein expressed in *E. coli*, immunizing mice with the protein, and producing a hybridoma cell by fusing the spleen cells and myeloma cells (Current Protocols in Molecular Biology (1987) Ausubel et al. edit, John Wily & Sons, Section 11.4-11.11).

[0057] The antibody binding to the protein of the present invention can be used for purification of the protein of the invention, and also for detection and/or diagnosis of the abnormalities of the expression and structure of the protein. Specifically, proteins can be extracted, for example, from tissues, blood, or cells, and the protein of the invention is detected by Western blotting, immunoprecipitation, or ELISA, etc. for the above purpose.

[0058] Furthermore, the antibody binding to the protein of the present invention can be utilized for treating the diseases that associates with the protein of the invention. If the antibodies are used for treating patients, human antibodies or humanized antibodies are preferable in terms of their low antigenicity. The human antibodies can be prepared by immunizing a mouse whose immune system is replaced with that of human ("Functional transplant of megabase human immunoglobulin loci recapitulates human antibody response in mice" Mendez M.J. et al. (1997) Nat. Genet. 15: 146-156). The humanized antibodies can be prepared by recombination of the hypervariable region of a monoclonal antibody (Methods in Enzymology (1991) 203: 99-121).

[0059] The cDNA of the present invention encodes the amino acid sequence of a protein which is predicted to have the function(s) described below based on the homology search of the GenBank and SwissProt. Specifically, for instance, as shown in EXAMPLES, searching a known gene or protein that is homologous to the partial sequence of the full-length cDNA of the invention (5602 clone) and referring the function of the gene and of the protein encoded by the gene make it possible to predict the function of the protein encoded by the cDNA of the invention. In this way, each of 1437 clones out of the 5602 full-length cDNA clones of the invention was predicted to encode a protein that was classified into one or more of the following categories.

Secretory or membrane protein (261 clones)

Glycoprotein-associated protein (113 clones)

Signal transduction-associated protein (148 clones)

Transcription-associated protein (233 clones)

Disease-associated protein (437 clones)

Enzyme or metabolism-associated protein (301 clones)

Cell division- or cell proliferation-associated protein (74 clones)

Cytoskeleton-associated protein (92 clones)

RNA synthesis-associated protein (280 clones)

Nuclear protein (352 clones)

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Protein synthesis- or transport-associated protein (112 clones)

Cellular defense-associated protein (23 clones)

Development- or growth-associated protein (23 clones)

[0060] It is also possible to predict the protein function by looking into the amino acid sequence for the motifs such

as the signal sequence, transmembrane region, nuclear translocation signal, glycosylation signal, phosphorylation site, Zinc finger motif, and SH3 domain. The programs, PSORT (Nakai K., and Kanehisa M. (1992) Genomics 14: 897-911), SOSUI (Hirokawa T. et al. (1998) Bioinformatics 14: 378-379) (Mitsui Information Developing Inc.), and MEMSAT (Jones D.T., Taylor W.R., and Thornton J.M. (1994) Biochemistry 33: 3038-3049) can be used to predict the existence of the signal sequence or transmembrane region. Alternatively, a partial amino acid sequence of the protein is fused with another protein such as GFP, the fusion protein is transfected into cultured cells, and the localization is analyzed to predict the function of the original protein.

[0061] Based on the determined nucleotide sequences of the full-length cDNAs obtained in the present invention, it is possible to predict more detailed functions of the proteins encoded by the cDNA clones, for example, by searching the databases such as GenBank, Swiss-Prot and UniGene for homologies of the cDNAs; or by searching the amino acid sequences deduced from the full-length cDNAs for signal sequences by using software programs such as PSORT, for transmembrane regions by using software programs such as SOSUI or for motifs by using software programs such as Pfam (http://www.expasy.ch/prosite/). As a matter of course, the functions are often predictable by using partial sequence information (preferably 300 nucleotides or more) instead of the full-length nucleotide sequences. However, the result of the prediction by using partial nucleotide sequence does not always agree with the result obtained by using full-length nucleotide sequence, and thus, it is needless to say that the prediction of function is preferably performed based on the full-length nucleotide sequences of the 4997 clones (see Example 18). The amino acid sequences deduced from the full-length nucleotide sequences were searched for functional domains by PSORT, SOSUI and Pfam. Prediction of functions of proteins encoded by the clones and the categorization thereof were performed based on these results obtained.

The following 798 clones were categorized into secretory and/or membrane proteins.

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HEMBA1000356,
                   HEMBA1000518.
                                  HEMBA1000531, HEMBA1000637,
                                                                HEMBA1000719,
                                                                                HEMBA1000817.
    HEMBA1000822,
                   HEMBA1000852,
                                  HEMBA1000870,
                                                 HEMBA1000991.
                                                                HEMBA1001052,
                                                                                HEMBA1001071.
    HEMBA1001085,
                   HEMBA1001286,
                                  HEMBA1001351.
                                                 HEMBA1001407,
                                                                 HEMBA1001446.
                                                                                HEMBA1001515.
    HEMBA1001557,
                   HEMBA1001569,
                                   HEMBA1001661,
                                                 HEMBA1001734,
                                                                 HEMBA1001746.
                                                                                HEMBA1001866.
    HEMBA1002125,
                   HEMBA1002150,
                                  HEMBA1002166,
                                                 HEMBA1002417,
                                                                 HEMBA1002462,
                                                                                HEMBA1002475.
    HEMBA1002477,
                   HEMBA1002486,
                                  HEMBA1002609,
                                                 HEMBA1002659,
                                                                HEMBA1002661.
                                                                                HEMBA1002780.
    HEMBA1002818,
                   HEMBA1002876,
                                  HEMBA1002921,
                                                 HEMBA1003071,
                                                                HEMBA1003077,
                                                                                HEMBA1003079.
30
    HEMBA1003086,
                   HEMBA1003096,
                                   HEMBA1003281,
                                                 HEMBA1003286,
                                                                HEMBA1003538,
                                                                                HEMBA1003711,
    HEMBA1003742,
                   HEMBA1003803,
                                  HEMBA1004055,
                                                 HEMBA1004143,
                                                                HEMBA1004146,
                                                                                HEMBA1004207,
    HEMBA1004341,
                   HEMBA1004461,
                                  HEMBA1004577.
                                                 HEMBA1004637,
                                                                HEMBA1004752,
                                                                                HEMBA1004756,
    HEMBA1004850,
                   HEMBA1004889,
                                  HEMBA1004923.
                                                 HEMBA1004930,
                                                                HEMBA1005029,
                                                                                HEMBA1005035,
    HEMBA1005050,
                   HEMBA1005552.
                                  HEMBA1005576,
                                                 HEMBA1005581,
                                                                HEMBA1005588,
                                                                                HEMBA1005616,
35
    HEMBA1005699.
                   HEMBA1005991.
                                  HEMBA1006036.
                                                 HEMBA1006038,
                                                                HEMBA1006067,
                                                                                HEMBA1006173,
                   HEMBA1006293,
    HEMBA1006198.
                                  HEMBA1006310,
                                                 HEMBA1006492,
                                                                HEMBA1006502,
                                                                                HEMBA1006583,
    HEMBA1006659.
                   HEMBA1006758,
                                  HEMBA1006789,
                                                 HEMBA1006921,
                                                                HEMBA1006926,
                                                                                HEMBA1006976,
    HEMBA1007203,
                   HEMBA1007301.
                                  HEMBB1000037,
                                                 HEMBB1000050,
                                                                HEMBB1000054.
                                                                                HEMBB1000175,
    HEMBB1000317.
                   HEMBB1000556,
                                  HEMBB1000593,
                                                 HEMBB1000631,
                                                                HEMBB1000763,
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    HEMBB1000915,
                   HEMBB1000975,
                                  HEMBB1001112,
                                                 HEMBB1001151.
                                                                HEMBB1001177,
                                                                                HEMBB1001302.
    HEMBB1001348,
                   HEMBB1001564,
                                  HEMBB1001630,
                                                 HEMBB1001871,
                                                                HEMBB1001872,
                                                                                HEMBB1001925.
    HEMBB1001962.
                   HEMBB1002042,
                                  HEMBB1002044.
                                                 HEMBB1002142,
                                                                HEMBB1002190.
                                                                                HEMBB1002193.
    HEMBB1002247,
                   HEMBB1002383,
                                  HEMBB1002387.
                                                 HEMBB1002550.
                                                                HEMBB1002600.
                                                                                HEMBB1002692,
    MAMMA1000045, MAMMA1000129,
                                  MAMMA1000133,
                                                 MAMMA1000277.
                                                                MAMMA1000278.
                                                                               MAMMA1000410.
45
    MAMMA1000416, MAMMA1000472,
                                  MAMMA1000672,
                                                 MAMMA1000684.
                                                                MAMMA1000714,
                                                                               MAMMA1000734.
    MAMMA1000778, MAMMA1000798,
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                                                 MAMMA1000859,
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                                                                MAMMA1000897.
    MAMMA1001008, MAMMA1001030,
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    MAMMA1001754,
                   MAMMA1001771,
                                  MAMMA1002009.
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                                                                               MAMMA1002461,
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    MAMMA1002844,
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    NT2RM1000080.
                   NT2RM1000092.
                                  NT2RM1000131,
                                                 NT2RM1000199,
                                                                 NT2RM1000257,
                                                                                NT2RM1000260,
    NT2RM1000355,
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                                                 NT2RM1000857,
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    NT2RM1001115,
                   NT2RM1001139,
                                  NT2RM2000259.
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                                                                NT2RM2000287.
                                                                                NT2RM2000395,
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NT2RM2000490.

NT2RM2000522,

NT2RM2000566,

NT2RM2000422,

NT2RM2000402,

NT2RM2000407,

	NT2RM2000581,	NT2RM2000609,	NT2RM2000821,	NT2RM2001370,	NT2RM2001393,	NT2RM2001499,
	NT2RM2001547,	NT2RM2001613,	NT2RM2001648,	NT2RM2001659,	NT2RM2001671,	NT2RM2001688,
	NT2RM2001698,	NT2RM2001718,	NT2RM2001753,	NT2RM2001760,	NT2RM2001785,	NT2RM2001930,
	NT2RM2001950,	NT2RM2001997,	NT2RM2001998,	NT2RM2002049,	NT2RM2002145,	NT2RM4000233,
5	NT2RM4000433,	NT2RM4000457,	NT2RM4000486,	NT2RM4000496,	NT2RM4000520,	NT2RM4000634,
	NT2RM4000674,	NT2RM4000700,	NT2RM4000764,	NT2RM4000778,	NT2RM4000795,	NT2RM4000820,
	NT2RM4000857,	NT2RM4001032,	NT2RM4001054,	NT2RM4001116,	NT2RM4001455,	NT2RM4001666,
	NT2RM4001810,	NT2RM4001813,	NT2RM4001930,	NT2RM4001987,	NT2RM4002054,	NT2RM4002073,
	NT2RM4002145,	NT2RM4002146,	NT2RM4002189,	NT2RM4002194,	NT2RM4002251,	NT2RM4002339.
10	NT2RM4002438,	NT2RM4002446,	NT2RM4002452,	NT2RM4002460,	NT2RM4002493,	NT2RM4002558.
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	NT2RP1000358,	NT2RP1000413,	NT2RP1000418,	NT2RP1000547.	NT2RP1000609,	NT2RP1000677.
	NT2RP1000767,	NT2RP1000782,	NT2RP1000856,	NT2RP1001113,	NT2RP1001247,	NT2RP1001286,
	NT2RP1001310,	NT2RP1001311,	NT2RP1001313,	NT2RP1001385,	NT2RP1001449,	NT2RP1001546,
15	NT2RP1001569,	NT2RP2000032,	NT2RP2000040,	NT2RP2000056,	NT2RP2000070,	NT2RP2000091,
	NT2RP2000114,	NT2RP2000120,	NT2RP2000173,	NT2RP2000175.	NT2RP2000195.	NT2RP2000257,
	NT2RP2000270,	NT2RP2000283,	NT2RP2000288,	NT2RP2000289,	NT2RP2000459,	NT2RP2000516,
	NT2RP2000660,	NT2RP2000842,	NT2RP2000892,	NT2RP2001081.	NT2RP2001268,	NT2RP2001295,
	NT2RP2001366.	NT2RP2001378.	NT2RP2001576.	NT2RP2001581,	NT2RP2001597,	
20	NT2RP2001947,	NT2RP2001991,	NT2RP2002025.	NT2RP2002066.	NT2RP2002078,	NT2RP2001613,
	NT2RP2002312,	NT2RP2002325,	NT2RP2002385.	NT2RP2002479.	NT2RP2002537,	NT2RP2002105,
	NT2RP2002701,	NT2RP2002740,	NT2RP2002857,	NT2RP2003125,	NT2RP2003297,	NT2RP2002643,
	NT2RP2003446,	NT2RP2003466.	NT2RP2003506,	NT2RP2003513,		NT2RP2003433,
	NT2RP2003760,	NT2RP2003777.	NT2RP2003781,	NT2RP2003013,	NT2RP2003629,	NT2RP2003668,
25	NT2RP2004270,	NT2RP2004300,	NT2RP2004392,	NT2RP2004655.	NT2RP2004142,	NT2RP2004194,
	NT2RP2004799,	NT2RP2004936.	NT2RP2004959,	NT2RP2005012.	NT2RP2004681,	NT2RP2004775,
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	NT2RP2005784,	NT2RP2005812,	NT2RP2006069,	NT2RP2006100,	NT2RP2005752,	NT2RP2005781,
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30	NT2RP3000134,	NT2RP3000207,	NT2RP3000333.	NT2RP3000341,	NT2RP3000092, NT2RP3000393,	NT2RP3000109,
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	NT2RP3001497,	NT2RP3001538,	NT2RP3001589,	NT2RP3001642,	NT2RP3001708,	NT2RP3001453,
35	NT2RP3001727,	NT2RP3001739,	NT2RP3001799,	NT2RP3001943,		NT2RP3001716,
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	NT2RP3003059.	NT2RP3003071,	NT2RP3003101.	NT2RP3002969,	NT2RP3002985,	NT2RP3003008,
40	NT2RP3003242,	NT2RP3003302,	NT2RP3003353.	N12RP3003145,	NT2RP3003197,	NT2RP3003203,
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45	NT2RP4000417,	NT2RP4000500,	NT2RP4000524,	NT2RP4000556.	NT2RP4000259,	NT2RP4000323,
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50	NT2RP4001379,	NT2RP4001498,	NT2RP4001547,	•	NT2RP4001372,	NT2RP4001373,
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55	OVARC1000889, OVARC1000924,		OVARC1000722,	OVARC1000751,	OVARC1000850,	OVARC1000890,
		OVARC1000936,	OVARC1000959,	OVARC1000984,	OVARC1000999,	OVARC1001034,
	OVARC1001055,	OVARC1001117,	OVARC1001129,	OVARC1001154,	OVARC1001329,	OVARC1001381,
	OVARC1001391,	OVARC1001453,	OVARC1001476,	OVARC1001506,	OVARC1001610,	OVARC1001702,
	OVARC1001703,	OVARC1001713,	OVARC1001745,	OVARC1001767,	OVARC1002127,	OVARC1002138,

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                                                                    PLACE1008643,
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                                    PLACE3000413,
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    THYRO1001287,
                    THYRO1001320,
                                    THYRO1001401,
                                                    THYRO1001534,
                                                                    THYRO1001537,
                                                                                    THYRO1001541,
     THYRO1001828,
                     Y79AA1000258,
                                     Y79AA1000420,
                                                    Y79AA1000469,
                                                                    Y79AA1000734,
                                                                                    Y79AA1000800.
     Y79AA1000976,
                    Y79AA1001023,
                                    Y79AA1001177.
                                                    Y79AA1001384,
                                                                    Y79AA1001394,
                                                                                    Y79AA1001603,
     Y79AA1001647,
                    Y79AA1001846,
                                    Y79AA1001874.
                                                    Y79AA1002139,
                                                                    Y79AA1002246.
                                                                                    Y79AA1002351,
    Y79AA1002399, Y79AA1002416, MAMMA1002498, NT2RM4002287
     [0062] The following 142 clones were categorized into glycoprotein-associated proteins.
     HEMBA1000156, HEMBA1000518, HEMBA1000852,
                                                   HEMBA1001071, HEMBA1001286,
                                                                                   HEMBA1001661,
     HEMBA1001734,
                    HEMBA1001866.
                                    HEMBA1003071.
                                                   HEMBA1003077,
                                                                   HEMBA1003281,
                                                                                   HEMBA1003538,
     HEMBA1003679,
                    HEMBA1003866.
                                    HEMBA1005576.
                                                    HEMBA1005581.
                                                                   HEMBA1005699,
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     HEMBA1006976.
                                                    HEMBB1000915,
                    HEMBA1007301.
                                    HEMBB1000317,
                                                                   HEMBB1001871,
                                                                                   HEMBB1001872,
     HEMBB1002193, MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA1001388,
                                                                                   MAMMA1002329,
     MAMMA1002428, MAMMA1002573,
                                    MAMMA1003150, NT2RM1000648, NT2RM1001115,
                                                                                   NT2RM2000260.
    NT2RM2000407.
                    NT2RM2000422,
                                    NT2RM2000490,
                                                    NT2RM2001499,
                                                                    NT2RM2001659.
                                                                                   NT2RM2001930.
     NT2RM4000820,
                    NT2RM4000857,
                                    NT2RM4001810,
                                                    NT2RM4001813,
                                                                    NT2RM4001987.
                                                                                   NT2RM4002145,
    NT2RM4002189,
                    NT2RM4002251,
                                    NT2RM4002460,
                                                    NT2RM4002558,
                                                                    NT2RP1000677,
                                                                                    NT2RP1000782.
     NT2RP1000856,
                                    NT2RP2000056,
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                                                    NT2RP2000070,
                                                                    NT2RP2001295.
                                                                                    NT2RP2001378.
     NT2RP2001597,
                    NT2RP2001991,
                                    NT2RP2002025,
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                                                                    NT2RP2002385,
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    NT2RP4000212,
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                                    NT2RP4000724,
                                                    NT2RP4000817,
                                                                    NT2RP4000925,
                                                                                    NT2RP4001150,
     NT2RP4001372,
                    NT2RP4001730,
                                    NT2RP4001822,
                                                    NT2RP4001823.
                                                                    NT2RP5003522.
                                                                                    OVARC1000091,
     OVARC1000288,
                    OVARC1000682,
                                    OVARC1001055.
                                                    OVARC1001506,
                                                                    OVARC1001713,
                                                                                   OVARC1002127,
     PLACE1000213, PLACE1000401, PLACE1002437, PLACE1002583,
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PLACE1002722,
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    PLACE1008643, PLACE1009094,
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    PLACE4000230.
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                                                  PLACE4000581,
                                                                 THYRO1000327,
                                                                                 THYRO1000756,
    THYRO1001287, Y79AA1001603, Y79AA1001874, MAMMA1002498
    [0063] The following 140 clones were categorized into signal transduction-associated proteins.
    HEMBA1000303, HEMBA1000369, HEMBA1000608, HEMBA1000657,
                                                                  HEMBA1000919.
                                                                                 HEMBA1001019,
    HEMBA1001174,
                   HEMBA1001822,
                                   HEMBA1001921,
                                                  HEMBA1002139,
                                                                  HEMBA1002212,
                                                                                 HEMBA1002341,
    HEMBA1002417, HEMBA1002768,
                                   HEMBA1003250.
                                                  HEMBA1003291,
                                                                  HEMBA1003645,
                                                                                 HEMBA1004286.
    HEMBA1005737, HEMBA1006130, HEMBA1006708, HEMBB1000083,
                                                                  HEMBB1000266.
                                                                                 HEMBB1000632,
    HEMBB1000781, HEMBB1000831, HEMBB1002193, MAMMA1000173, MAMMA1001038,
    MAMMA1001198, MAMMA1002842, MAMMA1003057, NT2RM1000702,
                                                                  NT2RM1000772.
                                                                                 NT2RM1001072,
    NT2RM2000030.
                   NT2RM2000469.
                                   NT2RM2000612.
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    NT2RM4000229.
                   NT2RM4000354.
                                   NT2RM4000611.
                                                  NT2RM4000798.
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    NT2RM4001629.
                   NT2RM4001758.
                                   NT2RM4002013.
                                                  NT2RM4002527,
                                                                  NT2RP1000018,
                                                                                 NT2RP1000701,
    NT2RP1001294.
                   NT2RP1001302.
                                   NT2RP2000668,
                                                  NT2RP2001440,
                                                                  NT2RP2001560,
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    NT2RP2002193.
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                                                                  NT2RP2003164,
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    NT2RP2004232,
                   NT2RP2004768,
                                   NT2RP2006071,
                                                  NT2RP2006534,
                                                                  NT2RP3000759,
                                                                                 NT2RP3000845,
    NT2RP3001646.
                   NT2RP3001857,
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                                                  NT2RP3002004.
                                                                  NT2RP3002785,
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    NT2RP3002988,
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                                   NT2RP3004189.
                                                  NT2RP3004544,
                                                                  NT2RP4000147.
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                                                  NT2RP4002408.
                                                                  NT2RP5003477.
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    OVARC1000437, OVARC1000556, OVARC1000649, OVARC1000945, OVARC1001200.
    OVARC1002182,
                    PLACE1000977,
                                   PLACE1001387.
                                                  PLACE1002493.
                                                                  PLACE1002591.
                                                                                 PLACE1003190.
    PLACE1003353,
                   PLACE1004128,
                                   PLACE1004302,
                                                  PLACE1004937.
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    PLACE1008244,
                    PLACE1008650,
                                   PLACE1009468.
                                                  PLACE1009596.
                                                                  PLACE1009708,
                                                                                 PLACE1009845.
    PLACE1010926, PLACE1011041, PLACE2000164, PLACE2000371, PLACE3000145, PLACE3000350, THYRO
30
    1000072, THYRO1000748, THYRO1001120, Y79AA1000328, Y79AA1002431, HEMBA1001247, NT2RM2001813,
    NT2RM4001454, NT2RP2005140, NT2RP2005293,
                                                  NT2RP3000487, NT2RP3003311,
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    PLACE1003723, PLACE1005327, PLACE3000124,
    [0064] The following 321 clones were categorized into transcription -associated proteins.
    HEMBA1000158, HEMBA1000201, HEMBA1000216, HEMBA1000555, HEMBA1000561.
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    HEMBA1001077, HEMBA1001137, HEMBA1001405,
                                                  HEMBA1001510,
                                                                 HEMBA1001635,
                                                                                 HEMBA1001804.
    HEMBA1001809, HEMBA1001819, HEMBA1001847,
                                                  HEMBA1001869,
                                                                 HEMBA1002035.
                                                                                 HEMBA1002092,
    HEMBA1002177, HEMBA1002770,
                                  HEMBA1002935,
                                                  HEMBA1003408.
                                                                 HEMBA1003545,
                                                                                 HEMBA1003568.
    HEMBA1003662, HEMBA1003684,
                                  HEMBA1003760.
                                                  HEMBA1003953,
                                                                 HEMBA1004097,
                                                                                 HEMBA1004321.
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    HEMBA1004353, HEMBA1004389,
                                  HEMBA1004479.
                                                                 HEMBA1004973.
                                                  HEMBA1004758,
                                                                                 HEMBA1005219.
    HEMBA1005359, HEMBA1005513,
                                   HEMBA1005528,
                                                  HEMBA1005548.
                                                                 HEMBA1005558.
                                                                                 HEMBA1005931.
    HEMBA1006158, HEMBA1006248.
                                   HEMBA1006278,
                                                  HEMBA1006283.
                                                                 HEMBA1006347.
                                                                                 HEMBA1006359.
    HEMBA1006559, HEMBA1006941,
                                   HEMBB1000789.
                                                  HEMBB1001011.
                                                                 HEMBB1001314.
                                                                                 HEMBB1001482.
    HEMBB1001673, HEMBB1001749,
                                   HEMBB1001839.
                                                  HEMBB1001908.
                                                                 HEMBB1002134.
                                                                                 HEMBB1002217,
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    HEMBB1002342, HEMBB1002607, MAMMA1000183.
                                                 MAMMA1000388.
                                                                                MAMMA1001222,
                                                                 MAMMA1001105,
    MAMMA1001260, MAMMA1001627, MAMMA1001633, MAMMA1001743, MAMMA1001820,
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    MAMMA1002617. MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000086,
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                                   NT2RM1001092,
    NT2RM1000885,
                   NT2RM1000894.
                                                  NT2RM2000013,
                                                                  NT2RM2000452,
                                                                                 NT2RM2000735.
    NT2RM2000740.
                   NT2RM2001035,
                                   NT2RM2001105,
                                                  NT2RM2001575,
                                                                  NT2RM2001670.
                                                                                 NT2RM2001716.
    NT2RM2001771,
                                   NT2RM4000024,
                   NT2RM2002091,
                                                  NT2RM4000046,
                                                                  NT2RM4000104,
                                                                                 NT2RM4000202.
    NT2RM4000531, NT2RM4000595, NT2RM4000733, NT2RM4000734,
    NT2RM4000741,
                   NT2RM4000751,
                                   NT2RM4000996,
                                                  NT2RM4001092,
                                                                  NT2RM4001140,
                                                                                 NT2RM4001200,
    NT2RM4001483,
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                                   NT2RM4001783,
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                                                   NT2RP1000111,
                                                                  NT2RP1000574,
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     [0065] The following 392 clones were categorized into disease-associated proteins.
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     [0066] The following 427 clones presumably belong to enzymes and/or metabolism-associated proteins.
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                                                                   HEMBB1001294,
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                    PLACE1010870,
                                    PLACE1011056.
                                                    PLACE1011433.
                                                                   PLACE1011664.
                                                                                   PLACE2000014,
     PLACE2000427,
                    PLACE3000009,
                                    PLACE3000169,
                                                   PLACE4000014.
                                                                   PLACE4000156.
                                                                                   PLACE4000192,
     PLACE4000261,
                    PLACE4000326,
                                   PLACE4000489,
                                                   SKNMC1000011,
                                                                   THYRO1000085.
                                                                                   THYRO1000242.
     THYRO1000585,
                    THYRO1001100,
                                    THYRO1001189,
                                                    THYRO1001809,
                                                                    Y79AA1000037,
                                                                                   Y79AA1000214,
     Y79AA1000231,
                    Y79AA1000589,
                                    Y79AA1000752.
                                                    Y79AA1001391,
                                                                    Y79AA1001613,
                                                                                    Y79AA1001705,
     Y79AA1001963, Y79AA1002431, Y79AA1002472, Y79AA1002482,
     [0069] The following 296 clones presumably belong to DNA- and/or RNA-binding proteins. HEMBA1000158,
     HEMBA1000216,
                    HEMBA1000561, HEMBA1000591, HEMBA1000851,
                                                                   HEMBA1001088,
                                                                                   HEMBA1001137,
     HEMBA1001405.
                    HEMBA1001510,
                                   HEMBA1001804,
                                                   HEMBA1001809,
                                                                   HEMBA1001819,
                                                                                   HEMBA1001847,
    HEMBA1001869.
                    HEMBA1002177,
                                    HEMBA1002935,
                                                   HEMBA1003408,
                                                                   HEMBA1003545,
                                                                                   HEMBA1003568,
     HEMBA1003591,
                    HEMBA1003662.
                                    HEMBA1003684,
                                                   HEMBA1003760,
                                                                   HEMBA1003783,
                                                                                   HEMBA1003805,
     HEMBA1003953.
                    HEMBA1004321,
                                   HEMBA1004354,
                                                   HEMBA1004389,
                                                                   HEMBA1004479,
                                                                                   HEMBA1004669,
     HEMBA1004847.
                    HEMBA1004973,
                                    HEMBA1005202,
                                                   HEMBA1005359,
                                                                   HEMBA1005931,
                                                                                   HEMBA1006248.
    HEMBA1006278,
                    HEMBA1006283,
                                   HEMBA1006359,
                                                   HEMBA1006652.
                                                                   HEMBA1007087,
                                                                                   HEMBA1007194.
    HEMBB1000264.
                   HEMBB1000789,
                                    HEMBB1001011,
                                                   HEMBB1001482,
                                                                   HEMBB1001736.
                                                                                   HEMBB1001749,
     HEMBB1001839, HEMBB1002217,
                                   MAMMA1000183.
                                                  MAMMA1000284.
                                                                  MAMMA1000731.
                                                                                  MAMMA1001105.
    MAMMA1001222, MAMMA1001260, MAMMA1001743, MAMMA1001837.
                                                                  MAMMA1002385, MAMMA1002617,
    MAMMA1002869, MAMMA1002937.
                                    MAMMA1003011, NT2RM1000086.
                                                                   NT2RM1000539.
                                                                                   NT2RM1000555,
    NT2RM1000666,
                    NT2RM1000691,
                                    NT2RM1000826.
                                                   NT2RM1000885.
                                                                   NT2RM1001059,
                                                                                   NT2RM1001092,
45
    NT2RM2000371,
                    NT2RM2000624,
                                    NT2RM2000735.
                                                   NT2RM2001105.
                                                                   NT2RM2001424,
                                                                                   NT2RM2001575,
    NT2RM2001605,
                    NT2RM2001670,
                                    NT2RM2001771,
                                                   NT2RM2001823.
                                                                                   NT2RM2002004,
                                                                   NT2RM2001989,
    NT2RM2002014,
                    NT2RM2002088,
                                    NT2RM2002091,
                                                   NT2RM4000046,
                                                                   NT2RM4000104,
                                                                                   NT2RM4000167.
    NT2RM4000191,
                    NT2RM4000202,
                                    NT2RM4000531,
                                                   NT2RM4000595.
                                                                   NT2RM4000733,
                                                                                   NT2RM4000751.
    NT2RM4000996, NT2RM4001092, NT2RM4001140,
    NT2RM4001178,
                    NT2RM4001200,
                                    NT2RM4001483,
                                                   NT2RM4001592,
                                                                   NT2RM4001783,
                                                                                   NT2RM4001823,
    NT2RM4001828.
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                                    NT2RM4001880,
                                                   NT2RM4001979,
                                                                   NT2RM4002093,
                                                                                   NT2RM4002109.
    NT2RP1000470,
                    NT2RP1000493,
                                    NT2RP1000574,
                                                   NT2RP1000902,
                                                                   NT2RP1000966,
                                                                                   NT2RP1001013,
    NT2RP1001073,
                    NT2RP1001080,
                                    NT2RP2000008,
                                                   NT2RP2000153,
                                                                   NT2RP2000258,
                                                                                   NT2RP2000297.
    NT2RP2001127,
                    NT2RP2001174,
                                    NT2RP2001233,
                                                   NT2RP2001511,
                                                                   NT2RP2001756,
                                                                                   NT2RP2001869,
    NT2RP2002079,
                    NT2RP2002099,
                                    NT2RP2002503,
                                                   NT2RP2002591,
                                                                   NT2RP2002939,
                                                                                   NT2RP2003157,
    NT2RP2003329,
                    NT2RP2003347,
                                    NT2RP2003480,
                                                   NT2RP2003522,
                                                                   NT2RP2003564,
                                                                                   NT2RP2003714,
    NT2RP2004187,
                    NT2RP2004568,
                                    NT2RP2004920.
                                                   NT2RP2005003.
                                                                   NT2RP2005139,
                                                                                   NT2RP2005168,
    NT2RP2005436,
                    NT2RP2005496,
                                    NT2RP2005701,
                                                   NT2RP2005763,
                                                                   NT2RP2005776,
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     NT2RP3000562,
                    NT2RP3000590,
                                     NT2RP3000603.
                                                     NT2RP3000624.
                                                                     NT2RP3000632.
                                                                                     NT2RP3000994.
     NT2RP3001057,
                     NT2RP3001107,
                                     NT2RP3001120,
                                                     NT2RP3001150,
                                                                     NT2RP3001155.
                                                                                     NT2RP3001338,
     NT2RP3001398,
                    NT2RP3001472,
                                     NT2RP3001672,
                                                     NT2RP3001688,
                                                                     NT2RP3001724,
                                                                                     NT2RP3001792,
     NT2RP3001855,
                    NT2RP3002165.
                                     NT2RP3002399,
                                                     NT2RP3002876,
                                                                     NT2RP3003138,
                                                                                     NT2RP3003193.
     NT2RP3003251,
                    NT2RP3003327,
                                     NT2RP3003555,
                                                     NT2RP3004013,
                                                                     NT2RP3004078.
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                                                     NT2RP3004617,
                                                                     NT2RP3004618,
                                                                                     NT2RP4000111.
     NT2RP4000398,
                    NT2RP4000455,
                                     NT2RP4000518,
                                                    NT2RP4000648,
                                                                     NT2RP4000865,
                                                                                     NT2RP4000929.
     NT2RP4001080,
                    NT2RP4001095,
                                     NT2RP4001213,
                                                    NT2RP4001433,
                                                                     NT2RP4001568,
                                                                                     NT2RP4001696.
     NT2RP4001753,
                    NT2RP4001838,
                                    NT2RP4001938,
                                                    NT2RP4002078,
                                                                    OVARC1000006,
                                                                                    OVARC1000087.
     OVARC1000241,
                    OVARC1000746,
                                     OVARC1000846,
                                                    OVARC1001232,
                                                                    OVARC1001271,
                                                                                    OVARC1001306,
     OVARC1001987,
                     OVARC1002112,
                                     PLACE1000406,
                                                     PLACE1000583,
                                                                     PLACE1000979,
                                                                                     PLACE1001118,
     PLACE1001632,
                    PLACE1001739,
                                     PLACE1002438,
                                                     PLACE1002532,
                                                                     PLACE1002775,
                                                                                     PLACE1002834,
     PLACE1003302.
                    PLACE1003519,
                                     PLACE1003605.
                                                     PLACE1003704.
                                                                     PLACE1003738,
                                                                                     PLACE1003885,
     PLACE1004471,
                    PLACE1004564,
                                     PLACE1004814,
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                                                                     PLACE1005876,
                                                                                     PLACE1005951,
     PLACE1006196,
                    PLACE1006482.
                                     PLACE1006488,
                                                     PLACE1006531,
                                                                     PLACE1006917,
                                                                                     PLACE1007346,
                    PLACE1007598.
                                     PLACE1007688,
     PLACE1007547.
                                                    PLACE1007969,
                                                                     PLACE1008132,
                                                                                     PLACE1009099,
     PLACE1009246.
                    PLACE1009398.
                                     PLACE1009476,
                                                    PLACE1009622,
                                                                     PLACE1010053,
                                                                                     PLACE1010194,
     PLACE1010702,
                    PLACE1010870,
                                     PLACE1011056,
                                                     PLACE1011114,
                                                                     PLACE1011433,
                                                                                     PLACE2000427,
     PLACE3000009.
                    PLACE3000169,
                                     PLACE4000014,
                                                     PLACE4000156,
                                                                     PLACE4000192,
                                                                                     PLACE4000261,
     PLACE4000489,
                    SKNMC1000091,
                                    THYRO 1000085,
                                                     THYRO1000242.
                                                                     THYRO1000501,
                                                                                    THYRO1001100.
     THYRO1001189,
                     THYRO1001809,
                                     Y79AA1000037.
                                                     Y79AA1000349,
                                                                     Y79AA1000752.
                                                                                     Y79AA1001211.
     Y79AA1001312,
                     Y79AA1001391,
                                     Y79AA1001613.
                                                     Y79AA1002103,
                                                                     Y79AA1002472.
                                                                                     Y79AA1002482,
     HEMBA1004596, OVARC1000148, PLACE1003334, THYRO1001661,
     [0070] The following 66 clones presumably belong to the category of RNA synthesis-associated proteins.
     HEMBA1000591,
                    HEMBA1001579,
                                    HEMBA1003179.
                                                    HEMBA1003591.
                                                                    HEMBA1006278.
                                                                                    HEMBB1000226,
     NT2RM1000187,
                    NT2RM1000852,
                                    NT2RM2000624.
                                                    NT2RM2001989.
                                                                    NT2RM2002100,
                                                                                    NT2RM4000191,
     NT2RM4001178,
                    NT2RM4002093,
                                     NT2RP1000035,
                                                    NT2RP1000272.
                                                                    NT2RP1000470,
                                                                                    NT2RP1001080,
     NT2RP2000153,
                    NT2RP2002928,
                                     NT2RP2003157,
                                                    NT2RP2004568,
                                                                    NT2RP2005126,
                                                                                    NT2RP2005436,
30
     NT2RP2005539,
                    NT2RP2005605,
                                     NT2RP2005776.
                                                    NT2RP2005942,
                                                                    NT2RP2006043,
                                                                                    NT2RP2006238.
     NT2RP3000361,
                    NT2RP3000397.
                                     NT2RP3001671,
                                                    NT2RP3004504,
                                                                     NT2RP4000078,
                                                                                     NT2RP4000111.
     NT2RP4000481,
                    NT2RP4000518,
                                    NT2RP4000614,
                                                    NT2RP4000929,
                                                                    NT2RP4001696,
                                                                                     NT2RP4002058,
     OVARC1001232.
                    OVARC1001577,
                                     PLACE1000406,
                                                    PLACE1000596,
                                                                    PLACE1000755,
                                                                                     PLACE1001739,
     PLACE1003704.
                    PLACE1003885,
                                     PLACE1004564,
                                                    PLACE1004814,
                                                                    PLACE1004902,
                                                                                     PLACE1005373,
     PLACE1005646.
                    PLACE1005876,
                                     PLACE1006196,
                                                    PLACE1006626,
                                                                    PLACE1006878,
                                                                                     PLACE1006917,
     PLACE1009476, PLACE1009925, PLACE1010194, PLACE1011114, THYRO1000121, Y79AA1001963,
    [0071] The following 184 clones presumably belong to protein synthesis- and/or protein transport-associated pro-
    teins.
     HEMBA1000012,
                    HEMBA1000141,
                                    HEMBA1000592,
                                                    HEMBA1003617,
                                                                    HEMBA1003773.
                                                                                    HEMBA1004202.
40
    HEMBA1004276,
                    HEMBA1004734,
                                    HEMBA1004847.
                                                    HEMBA1004929,
                                                                    HEMBA1004930.
                                                                                    HEMBA1005047.
    HEMBA1005202,
                    HEMBA1006031.
                                    HEMBA1006272,
                                                    HEMBA1006474.
                                                                    HEMBA1006652,
                                                                                    HEMBA1006914,
    HEMBA1006973,
                    HEMBA1007224.
                                    HEMBB1000915.
                                                    HEMBB1001112.
                                                                    HEMBB1001137,
                                                                                    HEMBB1001736,
    HEMBB1001831.
                   HEMBB1001915,
                                   MAMMA1000085, MAMMA1000734,
                                                                   MAMMA1001008.
                                                                                   MAMMA1002170,
    MAMMA1002219, MAMMA1002236.
                                    MAMMA1002619, NT2RM1000661, NT2RM1000833,
                                                                                    NT2RM2000092,
45
    NT2RM2000504,
                    NT2RM2000577.
                                    NT2RM2000821.
                                                    NT2RM2001201,
                                                                    NT2RM2001592,
                                                                                    NT2RM2001613,
    NT2RM2001648,
                    NT2RM2001730,
                                    NT2RM2001760.
                                                    NT2RM2002055,
                                                                    NT2RM4000155,
                                                                                    NT2RM4000169.
    NT2RM4000344,
                    NT2RM4000356.
                                    NT2RM4000421,
                                                    NT2RM4000712,
                                                                    NT2RM4001054,
                                                                                    NT2RM4001203.
    NT2RM4001382,
                    NT2RM4001444.
                                    NT2RM4002062,
                                                    NT2RM4002205,
                                                                    NT2RM4002623,
                                                                                    NT2RP1000326,
    NT2RP1000522.
                    NT2RP1000547,
                                    NT2RP1000746,
                                                    NT2RP1000947,
                                                                    NT2RP1001569.
                                                                                    NT2RP2000147,
    NT2RP2000710,
                    NT2RP2000880,
                                    NT2RP2000943,
                                                    NT2RP2001290,
                                                                    NT2RP2001392,
                                                                                     NT2RP2001601.
    NT2RP2001613,
                    NT2RP2001660,
                                    NT2RP2001740,
                                                    NT2RP2002124,
                                                                    NT2RP2002606.
                                                                                    NT2RP2002862.
    NT2RP2002959,
                    NT2RP2002980,
                                    NT2RP2003137,
                                                    NT2RP2003158,
                                                                    NT2RP2003391,
                                                                                    NT2RP2003394,
    NT2RP2003401,
                    NT2RP2003433,
                                    NT2RP2003704,
                                                    NT2RP2003713,
                                                                    NT2RP2003737,
                                                                                    NT2RP2003760,
    NT2RP2003981,
                    NT2RP2004366,
                                     NT2RP2004389,
                                                    NT2RP2004791,
                                                                    NT2RP2005012,
                                                                                     NT2RP2005116,
55
    NT2RP2005360, NT2RP2005763, NT2RP2005784, NT2RP3000366,
    NT2RP3000759,
                    NT2RP3000968,
                                     NT2RP3001113,
                                                    NT2RP3001690,
                                                                    NT2RP3002045,
                                                                                    NT2RP3002151,
    NT2RP3002529.
                    NT2RP3002671,
                                     NT2RP3003301.
                                                    NT2RP3003846,
                                                                    NT2RP3003876,
                                                                                    NT2RP3004209,
    NT2RP4000370,
                    NT2RP4000457,
                                     NT2RP4000879,
                                                    NT2RP4000927,
                                                                    NT2RP4001041,
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NT2RP4001313.
                    NT2RP4001315.
                                    NT2RP4001574.
                                                   NT2RP4001592.
                                                                   OVARC1000013,
                                                                                   OVARC1000071,
     OVARC1000085.
                    OVARC1000465.
                                    OVARC1000564,
                                                   OVARC1000771,
                                                                   OVARC1000862,
                                                                                   OVARC1001171,
     OVARC1001180,
                    OVARC1001342.
                                    PLACE1000007,
                                                    PLACE1000061,
                                                                   PLACE1000081,
                                                                                   PLACE1000492.
     PLACE1000863.
                    PLACE1001092,
                                    PLACE1001748,
                                                    PLACE1002090,
                                                                   PLACE1003174.
                                                                                   PLACE1003915,
    PLACE1004104,
                    PLACE1004270,
                                    PLACE1004743.
                                                    PLACE1005557.
                                                                   PLACE1005813,
                                                                                   PLACE1006170.
     PLACE1006488.
                    PLACE1006829.
                                    PLACE1007706,
                                                    PLACE1007729,
                                                                   PLACE1008273.
                                                                                   PLACE1008402.
     PLACE1008790,
                    PLACE1008813,
                                    PLACE1009094.
                                                    PLACE1009130.
                                                                   PLACE1009477.
                                                                                   PLACE1009721.
     PLACE1009845,
                    PLACE1010074.
                                    PLACE1010547.
                                                    PLACE1011109.
                                                                   PLACE1011229.
                                                                                   PLACE1011477,
     PLACE1012031,
                    PLACE2000404,
                                    PLACE3000059,
                                                    PLACE3000121,
                                                                   PLACE4000269.
                                                                                   PLACE4000654,
     SKNMC1000011,
                    THYRO1000983,
                                    THYRO1001003.
                                                    THYRO1001313,
                                                                    Y79AA1000560,
                                                                                   Y79AA1000784.
     Y79AA1000968, Y79AA1001493, Y79AA1001875, Y79AA1002027, Y79AA1002209, HEMBA1006284,
     [0072] The following 130 clones presumably belong to cytoskeleton-associated proteins.
     HEMBA1000156, HEMBA1000168, HEMBA1000411, HEMBA1000588, HEMBA1001043,
                                                                                  HEMBA1001651.
     HEMBA1001661.
                    HEMBA1002102,
                                    HEMBA1002161,
                                                   HEMBA1002939,
                                                                   HEMBA1003235,
                                                                                  HEMBA1003581,
    HEMBA1004499,
                    HEMBA1004534,
                                    HEMBA1004697,
                                                   HEMBA1004929,
                                                                   HEMBA1004972,
                                                                                  HEMBA1005582,
     HEMBA1005595,
                    HEMBA1006344,
                                    HEMBA1006737,
                                                   HEMBB1001175,
                                                                   HEMBB1001282,
                                                                                  HEMBB1001562,
     HEMBB1001802, MAMMA1000824, MAMMA1001041, MAMMA1001576,
                                                                   MAMMA1001679, MAMMA1001735,
     MAMMA1002297, MAMMA1002351, MAMMA1002622, MAMMA1002637, MAMMA1003127, NT2RM1000850,
     NT2RM1000898.
                    NT2RM2000030,
                                    NT2RM2000260,
                                                   NT2RM2000691,
                                                                   NT2RM2001324.
                                                                                  NT2RM4000169,
20
     NT2RM4000229, NT2RM4000515, NT2RM4001217, NT2RP1000202, NT2RP1000348, NT2RP1000460,
     NT2RP1000478,
                    NT2RP1001033.
                                    NT2RP1001294.
                                                   NT2RP1001302,
                                                                   NT2RP2000070.
                                                                                   NT2RP2000812,
     NT2RP2000814.
                    NT2RP2001168.
                                    NT2RP2001245.
                                                   NT2RP2001634.
                                                                   NT2RP2001900.
                                                                                   NT2RP2003307,
     NT2RP2003394.
                    NT2RP2004041.
                                    NT2RP2004242,
                                                   NT2RP2004538,
                                                                   NT2RP2004587,
                                                                                   NT2RP2004681,
     NT2RP2004732.
                    NT2RP2004978.
                                    NT2RP2005491.
                                                   NT2RP2005531,
                                                                   NT2RP2005712,
                                                                                   NT2RP2006275,
     NT2RP3000753.
                    NT2RP3001113.
                                    NT2RP3001216,
                                                   NT2RP3001239.
                                                                   NT2RP3001272,
                                                                                   NT2RP3001554,
     NT2RP3001690.
                    NT2RP3001799.
                                    NT2RP3002688.
                                                   NT2RP3003061,
                                                                   NT2RP3003185,
                                                                                   NT2RP3003230,
    NT2RP3004569,
                    NT2RP3004578,
                                    NT2RP4001004,
                                                   NT2RP4001086,
                                                                   NT2RP4001256,
                                                                                   NT2RP4001567.
    NT2RP4001927,
                    OVARC1000001.
                                    OVARC1000106,
                                                   OVARC1000437.
                                                                   OVARC1000520,
                                                                                   OVARC1000679,
     OVARC1001731, OVARC1002050, PLACE1001104, PLACE1002571,
30
     PLACE1002591,
                    PLACE1002655,
                                    PLACE1002714.
                                                   PLACE1003625.
                                                                   PLACE1005287,
                                                                                   PLACE1006552.
     PLACE1007946,
                    PLACE1008426.
                                    PLACE1010148.
                                                   PLACE1010547.
                                                                   PLACE1010743.
                                                                                   PLACE1010896.
     PLACE1010960,
                    PLACE1011310,
                                    PLACE1011922,
                                                   PLACE2000216,
                                                                   PLACE2000274.
                                                                                   PLACE2000371.
    PLACE2000458,
                    PLACE3000145,
                                    PLACE3000416,
                                                   PLACE4000009,
                                                                   THYRO1000132.
                                                                                   THYRO1001405.
     THYRO1001458, Y79AA1000368, Y79AA1000794, Y79AA1000833, Y79AA1000962, Y79AA1002208,
35
     [0073] The following 54 clones presumably belong to cell division-associated and/or cell proliferation-associated
     HEMBA1001019, HEMBA1001595,
                                                   HEMBA1002997, HEMBA1003136, HEMBA1003369,
                                    HEMBA1002363.
     HEMBA1004131, HEMBA1004354,
                                    HEMBA1005621.
                                                   HEMBB1000037, HEMBB1000264,
                                                                                  MAMMA1001768,
     MAMMA1002769. NT2RM1000354.
                                    NT2RM1000430,
                                                   NT2RM1000874,
                                                                   NT2RM2001256,
                                                                                  NT2RM2001743,
     NT2RM2001896.
                    NT2RM2002145,
                                    NT2RM4000215,
                                                    NT2RM4001714,
                                                                   NT2RP1000163,
                                                                                   NT2RP1000333,
     NT2RP1000439,
                    NT2RP2000346,
                                    NT2RP2001397,
                                                   NT2RP2002595,
                                                                   NT2RP2003177,
                                                                                   NT2RP2003596,
     NT2RP2003912.
                    NT2RP2004396,
                                    NT2RP2005037,
                                                   NT2RP2005520,
                                                                   NT2RP2005669,
                                                                                   NT2RP2005835,
     NT2RP3001730,
                    NT2RP3002081,
                                    NT2RP4000210,
                                                   NT2RP4000415,
                                                                   NT2RP4001414,
                                                                                   NT2RP4001634,
     OVARC1000013,
                    OVARC1000937,
                                    PLACE1001383,
                                                    PLACE1002433,
                                                                   PLACE1004316.
                                                                                   PLACE1005287,
     PLACE1008808, PLACE1010720, PLACE1010833, Y79AA1000748, Y79AA1001236, Y79AA1001394
     [0074] The following 36 clones presumably belong to the category of embryogenesis- and/or development-associ-
     ated proteins:
     HEMBA1000518,
                    HEMBA1001847, HEMBA1001869,
                                                   HEMBA1003545.
                                                                   HEMBA1004973.
                                                                                   HEMBB1002442.
     MAMMA1001837.
                    NT2RM2001670,
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     [0075] The following 30 clones presumably belong to cellular defense-associated proteins.
                                    HEMBA1003417, HEMBA1006253,
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NT2RP4001638,

PLACE1006958,

PLACE1008275,

NT2RP4001555,

NT2RP3002785,

NT2RP3004262,

PLACE1009113, PLACE1011858, PLACE4000014, THYRO1000684, Y79AA1002139, Y79AA1002229

[0076] Although it is unclear whether or not 261 clones out of clones other than the above-mentioned clones belong to any of the above-described categories, these clones are predicted to have some functions, based on the homology search using their full-length sequences.

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[0077] In some cases, the predicted functions based on the partial sequences are different from those based on the full-length sequences. The reason is that a protein does not always belong solely to a single category of the above-described functional categories, and therefore, a protein may belong to two or more of the predicted functional categories. Besides, additional functions can be found for the clones classified into these functional categories by further analyses.

[0078] Since the protein encoded by clones of the invention contains full-length amino acid sequence, it is possible to analyze its biological activity, and its effect on cellular conditions such as cell proliferation and differentiation by expressing the protein as a recombinant protein using an appropriate expression system, injecting the recombinant into the cell, or raising a specific antibody against the protein.

[0079] If the protein is a secretory protein, membrane protein, or protein associated with glycoprotein, signal trans-

duction, or transcription, its biological activity can be analyzed by the methods in "Gene Transcription" (Hames B.D., and Higgins S.J. edit, (1993)), "Glycobiology" (Fukuda M., and Kobata A. edit, (1993)), "Growth Factors" (McKay I., and Leigh I. edit, (1993)), "Extracellular Matrix" (Haralson M.A., and Hassell J.R. edit, (1995), "Transcription Factors" (Latchman D.S. edit, (1993)), "Signal Transduction" (Milligans G. edit, (1992)), "Protein Phosphorylation" (Hardies G. D. edit, (1993), and "Ion Channels" (Ashley R.H. edit, (1995) featured in "The Practical Approach Series" (IRL PRESS), or "Signal Transduction Protocols" (Kendall D.A., and Hill S.J. edit, (1995), "Glycoprotein Analysis in Biomedicine" (Hounsell E.F. edit, (1993)), featured in "Method in Molecular Biology" (Humana Press).

[0080] As to a protein associated with a disease, it is possible to perform a functional analysis as described above, but also possible to analyze correlation between the expression or the activity of the protein and a certain disease by using a specific antibody that is obtained by using expressed protein. Alternatively, it is possible to utilize the database Online Mendelian Inheritance in Man (OMIM) (http://www.ncbi.nlm.nih.gov/Omim/), which is a database of human genes and diseases, to analyze the protein.

New information is constantly being deposited in the OMIM database. Therefore, it is possible for one skilled in the art to find a new relationship between a particular disease and a gene of the present invention in the most up-to-date database.

[0081] Also, as for a secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., search of the OMIM with the following keywords resulted in the finding that the proteins are associated with many diseases (the result of the OMIM search for secrete and membrane proteins is shown below). Also, association between proteins associated to signal transduction or transcription and diseases is reported in "Transcription Factor Research-1999" (Fujii, Tamura, Kageyama, and Satake edit, (1999) Jikken-Igaku Zoukan, Vol.17, No.3), and "Gene Medicine" ((1999) Vol.3, No.2). For example, in tumors, many proteins have been shown to play a role, including secretory proteins, membrane proteins, and proteins associated with signal transduction, glycoprotein, and transcription, and also proteins associated with metabolism, cytoskeleton, and cell cycle, as described in "Tumor Biology" (Matsubara S. (1992) Syoukabou Life Science series). Thus, besides the proteins associated with diseases, many proteins described above are also potentially associated with diseases, and thus useful as a target in the medicinal industry.

[0082] The result of the OMIM search for secretory and membrane proteins is shown below, in which the keywords,

- (1) secretion protein,
- (2) membrane protein,
- (3) channel, and

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(4) extracellular matrix were used.

1017 entries found, searching for "membrane protein"

[0083] Shown in the search result are only the accession numbers in the OMIM. Using the number, data showing the relationship between a disease and a gene or protein can be seen. The OMIM data has been renewed everyday.

1) Secretion protein 268 entries found, searching for "secretion protein" 104760, 176860, 160900, 107400, 118910, 139320, 603850, 147572,176880, 600946, 603215, 157147, 600174, 40 151675, 170280, 179512, 179513,138120, 179509, 246700, 179510, 600626, 179511, 600998, 109270, 601489, 154545, 179490, 185860, 603216, 122559, 601746, 147290, 602672, 146770, 603062, 179508, 131230, 601591, 602421, 139250, 167805, 167770, 600041, 600564, 118825, 601146, 300090, 600753, 601652, 600759, 600768, 602434, 182590, 603166, 308230, 602534, 603489, 107470, 150390, 104610, 173120, 158106, 143890, 306900, 308700, 134797, 137350, 227500, 176300, 107730, 600760, 138079, 120180, 120160, 120150, 124092, 138160, 45 101000, 227600, 600509, 601199, 142410, 104311, 193400, 201910, 107300, 122560, 272800, 217000, 590050, 147670, 133170, 176730, 300300, 134370, 274600, 120140,162151, 158070, 152790, 120120, 106100, 300200, 192340, 190160, 138040, 147470, 147620, 173350, 147380, 152200, 152760, 157145, 153450, 264080, 113811, 600937, 600840, 188545, 202110, 600514, 186590, 603372, 136435, 137241, 252800, 214500, 207750, 138850, 139191, 142640, 138130, 189907, 603692, 600633, 603355, 107270, 600377, 147892, 232200, 600281, 232800, 50 602358, 137035, 601771, 601769, 253200, 601933, 118444, 600270, 120700, 600945, 603732, 147660, 600761, 172400, 600823, 600877, 130080, 171060, 107740, 307800, 602843, 130660, 152780, 124020, 601124, 601340, 601604, 601610, 171050, 312060, 232700, 300159, 142703, 600734, 125255, 168450, 123812, 188540, 147940, 188450, 600839, 182452, 188400, 182280, 176760, 263200, 600264, 188826, 252650, 601185, 162641, 137216, 601398, 601538, 118888, 118445, 601745, 190180, 601922, 182098, 602008, 147440, 602384, 600031, 109160, 55 602663, 151670, 602682, 602730, 602779, 146880, 603061, 142704, 603140, 106150, 600732, 153620, 603318, 139392, 600042, 102200, 603493, 182100, 264300, 603795, 184600 2) Membrane protein

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         3) Channel (member of membrane protein)
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         600235, 602023, 176263, 600681, 176265, 193245, 603305, 176258, 602983, 601219, 601141, 176267, 602343,
         602726, 138253, 176262, 600003, 600397, 602872, 138249, 600843, 600935, 600580, 600845, 602158, 602106,
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         304040, 154276, 180902, 121014, 602368, 139311, 601383, 108745, 601313, 601042, 600131, 186360, 600109,
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         600229, 600170, 603319, 601485, 118503, 180903, 602076, 124030, 601059, 601212, 601218, 147450, 600855,
         600919, 601154, 601157, 171060, 600968, 182139, 131230, 121015, 600421, 113730, 249210, 310500, 600637,
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         4) Extracellular matrix
         167 entries found, searching for "extracellular matrix"
         603479, 602201, 601418, 601548, 154870, 115437, 602285, 602262, 602261, 134797, 600754, 120361, 116935,
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         600261, 602109, 120140, 120150, 147557, 193400, 600536, 188826, 120180, 118661, 120320, 152200, 135821,
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         601652, 601692, 601728, 120220, 601915, 602048, 155120, 310300, 120210, 120165, 120120, 118940, 116930,
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         107269, 216550, 103320, 603489, 603551, 603767, 603799, 603842
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[0084] There are several methods for analyzing the expression levels of genes associated with diseases. Differences in gene expression levels between diseased and normal tissues are studied by the analytical methods, for example, Northern hybridization and differential display. Other examples include a method with high-density cDNA filter, a method with DNA microarray and methods with PCR amplification (Experimental Medicine, Vol.17, No. 8, 980-1056 (1999); Cell Engineering (additional volume) DNA Microarray and Advanced PCR Methods, Muramatsu & Naba (eds.), Shujunsya). The varying levels of gene expression between diseased tussues and normal tissues can be studied by any of these analytical methods. When explicit difference in the expression level is observed for a gene, it can be conclude that the gene is closely associated with a disease or disorder. Instead of diseased tissues, cultured cells can be used

for the assessment. Similarly, when gene expression is explicitly different between normal cells and cells reproducing disease-associated specific features, it can be concluded that the gene is closely associated with a disease or disorder. When the expression levels of genes are evidently varied during major cellular events (such as differentiation and apoptosis), the genes are involved in the cellular events and accordingly are candidates for disease- and/or disorder-associated genes. Further, genes exhibiting tissue-specific expression are genes playing important parts in the tissue functions and, therefore, can be candidates for genes associated with diseases and/or disorders affecting the tissues. [0085] For example, non-enzymic protein glycation reaction is believed to be a cause for a variety of chronic diabetic complications. Accordingly, genes of which expression levels are elevated or decreased in a glycated protein-dependent manner in the endothelial cells, are associated with diabetic complications caused by glycated proteins (Diabetes 1996, 45 (Suppl. 3), S67-S72; Diabetes, 1997, 46 (Suppl. 2), S19-S25).

The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)- α participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis. Many genes acting at the downstream of TNF- α and IL-1 β among inflammation-associated cytokines have been previously identified. The respective stimulations are transduced through independent pathways of signaling cascade. There exists another signaling cascade for both stimulations, wherein NF- κ B is a common transducing molecule shared by the two stimulations (J. Leukoc. Biol., 1994, 56(5): 542-547). It has also been revealed that many inflammation-associated genes, including IL-2, IL-6 and G-CSF, are varied in the expression levels thereof in response to the signal through the common pathway (Trend Genet. 1999, 15(6): 229-235). It is assumed that genes of which expression levels are varied in response to the stimulation of TNF- α or IL-1 β also participate in inflammation.

[0086] Ultraviolet radiation damage has been recognized as a risk factor for skin cancers, etc. (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with ultraviolet radiation damage of skin. In addition, genes associated with neural differentiation can be candidates for genes responsible for neurological diseases as well as candidates for genes usable for treating the diseases.

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[0087] Clones exhibiting differences in the expression levels thereof can be selected by using gene expression analysis. The selection comprises, for example; analyzing cDNA clones by using high-density cDNA filter; and statistically treating the multiple signal values (signal values of radioisotope in the labeled probes or values obtained by measuring fluorescence intensities emitted from the fluorescent labels) for the respective clones by two-sample t-test, where the signal values are determined by multiple experiments of hybridization. The clones of interest are selectable based on the statistically significant differences in the signal distribution at p<0.05. However, selectable clones with significant difference in the expression levels thereof may be changed depending on the partial modification of statistical treatment. For example, the clones may be selected by conducting statistical treatment with two-sample t-test at p<0.01; or genes exhibiting more explicit differences in the expression levels thereof can be selected by performing statistical treatment with a pre-determined cut-off value for the significant signal difference. An alternative method is that the expression levels are simply compared with each other, and then, the clones of interest are selected based on the ratio of the expression levels thereof.

[0088] Clones that vary in their expression levels can also be selected by comparing the expression levels by PCR analysis, for example, by using the method of determining the band intensities representing the amounts of PCR products with ethidium bromide staining; the method of determining the values of radioisotope signals or fluorescence intensities of the PCR products when radiolabeled or fluorescent dye-labeled primers, respectively, are used in PCR amplification; or the method of determining the values of radioisotope signals or fluorescence intensities of the probes hybridized to the PCR products when radiolabeled or fluorescent dye-labeled probes, respectively, are used in the hybridization. If the expression level ratios obtained in multiple PCR experiments are constantly at least 2-fold, such a clone can be judged to vary in its expression level. When the ratios are several-fold or not less than 10-fold, the clone can be selected as a gene exhibiting the explicit difference in its expression level.

[0089] A survey of genes of which expression levels are varied specifically to the glycated protein in the endothelial cells has revealed genes with elevated expression levels, HEMBA1003958, HEMBA1004850, MAMMA1001256, MAMMA1002132, PLACE2000411 and PLACE3000119. On the other hand, a gene of which expression level is decreased specifically to the glycated protein is MAMMA1001783. These clones are genes associated with diabetes.

[0090] A survey of genes of which expression levels are varied in response to TNF- α (Tumor Necrosis Factor-alpha) in the primary cell culture of synovial tissue has revealed the following clones with elevated expression levels in the presence of TNF- α :

HEMBA100005, HEMBA100012, HEMBA100020, HEMBA1000046, HEMBA100076, HEMBA1000111, HEMBA1000168, HEMBA1000185, HEMBA1000201, HEMBA1000231, HEMBA1000243, HEMBA1000280,

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HEMBA1000282, HEMBA1000304.
                                  HEMBA1000307. HEMBA1000327.
                                                                HEMBA1000356, HEMBA1000376,
    HEMBA1000387. HEMBA1000390.
                                  HEMBA1000418,
                                                 HEMBA1000460.
                                                                HEMBA1000491,
                                                                               HEMBA1000501.
    HEMBA1000518, HEMBA1000519,
                                  HEMBA1000520,
                                                 HEMBA1000531,
                                                                HEMBA1000534.
                                                                               HEMBA1000542,
    HEMBA1000545, HEMBA1000591,
                                  HEMBA1000592,
                                                 HEMBA1000594,
                                                                HEMBA1000636.
                                                                               HEMBA1000655.
    HEMBA1000657, HEMBA1000673,
                                  HEMBA1000682.
                                                 HEMBA1000686,
                                                                HEMBA1000722,
                                                                               HEMBA1000726.
    HEMBA1000827.
                   HEMBA1000870.
                                  HEMBA1000918,
                                                 HEMBA1000971,
                                                                HEMBA1000974.
                                                                               HEMBA1000986.
    HEMBA1001019,
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                                                 HEMBA1001059.
                                                                HEMBA1001060.
                                                                               HEMBA1001071.
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                                                 HEMBA1001172.
                                                                HEMBA1001196.
                                                                               HEMBA1001213,
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                                  HEMBA1001299,
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                                                                HEMBA1001303,
                                                                               HEMBA1001323.
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                   HEMBA1001327,
                                  HEMBA1001330.
                                                 HEMBA1001351,
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                                                 HEMBA1001912,
                                                                HEMBA1001921,
                                                                               HEMBA1001967.
    HEMBA1002084,
                   HEMBA1002161,
                                  HEMBA1002166,
                                                 HEMBA1002241,
                                                                HEMBA1002337.
                                                                               HEMBA1002363,
    HEMBA1002389,
                   HEMBA1002458,
                                  HEMBA1002460,
                                                 HEMBA1002469,
                                                                HEMBA1002538,
                                                                               HEMBA1002542,
    HEMBA1002547, HEMBA1002609, HEMBA1002624,
    HEMBA1002659,
                   HEMBA1002750,
                                  HEMBA1002770,
                                                 HEMBA1002779,
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                   HEMBA1002850,
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                                  HEMBA1003621.
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                                     NT2RP2005037,
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                                     NT2RP3001147,
                                                    NT2RP3001449,
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                                     NT2RP3001969,
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                                     OVARC1000971.
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                                                                                    OVARC1001085.
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                                                                    OVARC1001745.
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                                                     OVARC1002138.
                                                                                    PLACE1000005,
     PLACE1000420.
                     PLACE1000547.
                                     PLACE1000562,
                                                     PLACE1000653.
                                                                     PLACE1001168,
                                                                                     PLACE1001311,
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                     PLACE1001920,
                                     PLACE1001983.
                                                    PLACE1002066.
                                                                     PLACE1002072.
                                                                                    PLACE1002140,
     PLACE1002171,
                     PLACE1002319,
                                     PLACE1002474,
                                                    PLACE1002499.
                                                                     PLACE1002532,
                                                                                    PLACE1002665,
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                     PLACE1003145,
                                     PLACE1003361,
                                                    PLACE1003605.
                                                                     PLACE1003704,
                                                                                    PLACE1003783,
35
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                                     PLACE1004629,
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                                                                                    PLACE1005066,
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                     PLACE1005630,
                                     PLACE1005876,
                                                    PLACE1006143,
                                                                     PLACE1006325,
                                                                                    PLACE1006488,
     PLACE1006805,
                     PLACE1006829,
                                     PLACE1007286,
                                                    PLACE1007858,
                                                                     PLACE1008201,
                                                                                    PLACE1009045.
     PLACE1009113.
                     PLACE1009621,
                                     PLACE1010106,
                                                    PLACE1010310,
                                                                     PLACE1010622,
                                                                                    PLACE1010944,
     PLACE1010965.
                     PLACE1011185,
                                     PLACE1011332,
                                                    PLACE1011635,
                                                                     PLACE1011646,
                                                                                     PLACE1011725,
    PLACE2000014,
                     PLACE2000264,
                                     PLACE2000394,
                                                    PLACE2000419,
                                                                     PLACE3000160,
                                                                                    PLACE3000220,
    PLACE3000254,
                     PLACE3000271,
                                     PLACE3000339,
                                                    PLACE3000341,
                                                                     PLACE3000350,
                                                                                    PLACE3000353,
    PLACE3000401,
                    PLACE4000300,
                                    SKNMC1000091,
                                                    THYRO1000855,
                                                                     THYRO1001559,
                                                                                     Y79AA1000065,
     Y79AA1000202,
                     Y79AA1000214,
                                     Y79AA1000346,
                                                     Y79AA1000784,
                                                                     Y79AA1000833,
                                                                                     Y79AA1000968,
     Y79AA1001555, Y79AA1002220
     [0091] On the other hand, clones with decreased expression levels in the presence of TNF\alpha are:
    HEMBA1002150,
                     HEMBB1000240,
                                     NT2RM2000469,
                                                    NT2RM2000984,
                                                                    NT2RM2001688.
                                                                                    NT2RM4000290
    NT2RM4000496,
                     NT2RM4000590,
                                     NT2RM4001047,
                                                    NT2RM4001582.
                                                                    NT2RM4001611,
                                                                                    NT2RM4001650,
    NT2RM4002075.
                     NT2RM4002128,
                                     NT2RP1000174.
                                                     NT2RP1000243.
                                                                     NT2RP1000581,
                                                                                    NT2RP1000688,
    NT2RP1000767,
                     NT2RP1000825,
                                     NT2RP1001185:
                                                    NT2RP1001286,
                                                                    NT2RP1001432,
                                                                                    NT2RP1001457,
    NT2RP2000001,
                     NT2RP2000248.
                                     NT2RP2000841.
                                                    NT2RP2001813.
                                                                    NT2RP2002137,
                                                                                    NT2RP2002928,
    NT2RP2003517.
                     NT2RP2003559.
                                     NT2RP2003564.
                                                    NT2RP2004933,
                                                                    NT2RP2005038,
                                                                                    NT2RP2006365,
    NT2RP3000072,
                     NT2RP3000320,
                                     NT2RP3000484.
                                                     NT2RP3000980,
                                                                     NT2RP3001111,
                                                                                    NT2RP3001420,
                     NT2RP3002056,
    NT2RP3001495,
                                     NT2RP3002057,
                                                    NT2RP3002545,
                                                                    NT2RP3002713,
                                                                                    NT2RP3002799.
    NT2RP3002869.
                     NT2RP3002953,
                                     NT2RP3002955,
                                                    NT2RP3003282,
                                                                    NT2RP3003290.
                                                                                    NT2RP3003384.
    NT2RP3003385,
                     NT2RP3003870,
                                     NT2RP3004207,
                                                    NT2RP3004262,
                                                                    NT2RP3004527.
                                                                                    NT2RP4000500,
    NT2RP4000524,
                     NT2RP4000787,
                                     NT2RP4000927,
                                                    NT2RP4000955,
                                                                     NT2RP4000989,
                                                                                    NT2RP4001442.
    NT2RP4001638,
                     NT2RP4001950,
                                    NT2RP4002888,
                                                    NT2RP5003524,
                                                                    OVARC1001270.
                                                                                    PLACE1000246.
    PLACE1002816.
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These are rheumatoid arthritis-associated clones. A survey of genes of which expression levels are varied in primary cultured skin fibroblast cells exposed to ultraviolet light has revealed the following clones with elevated expression levels by ultraviolet radiation: HEMBA1000542, HEMBA1001808, HEMBA1002177, HEMBA1003314, MAMMA1001874. NT2RM2001100. NT2RP2005732, NT2RP3000592, NT2RP4000657, OVARC 1000004. OVARC1001092, OVARC1001342. PLACE1002816. NT2RM4001002. NT2RM4001813. NT2RM4002266, NT2RP2001174. NT2RP2001196. NT2RP2005358, NT2RP3000690, NT2RP3001216, NT2RP3003464, PLACE1006382. THYRO1000070. THYRO1001100, Y79AA1000342 [0094] On the other hand, the expression levels of the following clones were decreased by ultraviolet radiation: HEMBA1000005, HEMBA1000150, HEMBA1000156, HEMBA1000158, HEMBA1000168, HEMBA1000231. HEMBA1000304, HEMBA1000307. HEMBA1000333, HEMBA1000366, HEMBA1000369. HEMBA1000390, HEMBA1000396, HEMBA1000418, HEMBA1000434, HEMBA1000464, HEMBA1000469, HEMBA1000490. HEMBA1000504, HEMBA1000505, HEMBA1000557, HEMBA1000657, HEMBA1000673. HEMBA1000682, HEMBA1000686, HEMBA1000727, HEMBA1000752, HEMBA1000851, HEMBA1000852, HEMBA1000870, HEMBA1000872, HEMBA1001085, HEMBA1001121, HEMBA1001133, HEMBA1001235, HEMBA1001265, HEMBA1001281, HEMBA1001289, HEMBA1001299, HEMBA1001303, HEMBA1001310, HEMBA1001323, HEMBA1001595, HEMBA1001620, HEMBA1001640, HEMBA1001678, HEMBA1001712, HEMBA1001835, HEMBA1001950, HEMBA1001987. HEMBA1002253, HEMBA1002321, HEMBA1002341, HEMBA1002419. HEMBA1002679, HEMBA1002728, HEMBA1002818, HEMBA1002935, HEMBA1002999. HEMBA1003034, 20 HEMBA1003071, HEMBA1003098, HEMBA1003142, HEMBA1003175. HEMBA1003202, HEMBA1003212, HEMBA1003220, HEMBA1003276, HEMBA1003373, HEMBA1003417. HEMBA1003447, HEMBA1003528, HEMBA1003684, HEMBA1003799. HEMBA1003885. HEMBA1003989, HEMBA1004011, HEMBA1004055, HEMBA1004133. HEMBA1004225. HEMBA1004272, HEMBA1004353. HEMBA1004631, HEMBA1004669, HEMBA1004705. HEMBA1004753. HEMBA1004776, HEMBA1004803, HEMBA1004816, HEMBA1004900, HEMBA1005047, HEMBA1005079. HEMBA1005101, HEMBA1005149, HEMBA1005152, HEMBA1005202, HEMBA1005314, HEMBA1005372. HEMBA1005511, HEMBA1005513, HEMBA1005518, HEMBA1005570, HEMBA1005588, HEMBA1005577, HEMBA1005581, HEMBA1005609, HEMBA1005632, HEMBA1005853. HEMBA1006031, HEMBA1006035, HEMBA1006485, HEMBA1006486, HEMBA1006502. HEMBA1006696, HEMBA1006789, HEMBA1006796, HEMBA1007085, HEMBA1007224, HEMBA1007301. HEMBA1007319, 30 HEMBA1007341, HEMBA1007342, HEMBB1000036, HEMBB1000037. HEMBB1000217. HEMBB1000266, HEMBB1000317, HEMBB1000336, HEMBB1000354, HEMBB1000369. HEMBB1000399, HEMBB1000434. HEMBB1000438, HEMBB1000592, HEMBB1000673, HEMBB1000789, HEMBB1000810. HEMBB1000883. HEMBB1000887, HEMBB1001105, HEMBB1001182, HEMBB1001242, HEMBB1001267, HEMBB1001424, HEMBB1001464, HEMBB1001531, HEMBB1001618, HEMBB1001996, HEMBB1002092, HEMBB1002139. 35 HEMBB1002142, HEMBB1002190, HEMBB1002453, HEMBB1002520, HEMBB1002550. HEMBB1002556, HEMBB1002600, HEMBB1002664, MAMMA1000009, MAMMA1000055, MAMMA1000069. MAMMA1000133, MAMMA1000171, MAMMA1000173, MAMMA1000287, MAMMA1000416. MAMMA1000585, MAMMA1000713, MAMMA1000760, MAMMA1000798, MAMMA1000831. MAMMA1000875. MAMMA1000877, MAMMA1000876, MAMMA1000906, MAMMA1000931, MAMMA1000962. MAMMA1001139, MAMMA1001133, MAMMA1001243, MAMMA1001271, MAMMA1001274, MAMMA1001298, MAMMA1001606, MAMMA1001630, MAMMA1001670, MAMMA1001743, MAMMA1001751. MAMMA1002140. MAMMA1002145, MAMMA1002158, MAMMA1002170, MAMMA1002236, MAMMA1002311, MAMMA1002498. MAMMA1002754, MAMMA1002780, MAMMA1002820, MAMMA1002843, MAMMA1002844, MAMMA1002871, MAMMA1003047, NT2RM1000037, NT2RM1000039, NT2RM1000080. NT2RM1000086. NT2RM1000341, NT2RM1000499, NT2RM1000669, NT2RM1000746, NT2RM1000781. NT2RM1000885, NT2RM1000905, NT2RM1000962, NT2RM2000239, NT2RM2000260, NT2RM2000371, NT2RM2000639, NT2RM2000649, NT2RM2000735, NT2RM2000821, NT2RM2000984, NT2RM2001035, NT2RM2001065, NT2RM2001105, NT2RM2001177, NT2RM2001194, NT2RM2001196, NT2RM2001243, NT2RM2001256, NT2RM2001424. NT2RM2001588, NT2RM2001635, NT2RM2001648, NT2RM2001652, NT2RM2001668, NT2RM2001706, NT2RM2001727, NT2RM2001730. NT2RM2001743. NT2RM2001753, NT2RM2001760, NT2RM2001771, NT2RM2001785. NT2RM2001800. NT2RM2001855. NT2RM2001896, NT2RM2001997, NT2RM2002030, NT2RM2002049. NT2RM2002091, NT2RM2002142, NT2RM2002145, NT2RM2002178, NT2RM2002580, NT2RM4000215. NT2RM4000344. NT2RM4000368. NT2RM4000421, NT2RM4000425, NT2RM4000457. NT2RM4000496, NT2RM4000515, NT2RM4000712, NT2RM4000787, NT2RM4000813, NT2RM4000820. NT2RM4000852, NT2RM4000950. NT2RM4000996. NT2RM4001016, NT2RM4001047, NT2RM4001054, NT2RM4001140, NT2RM4001151, NT2RM4001187, NT2RM4001204, NT2RM4001258. NT2RM4001437, NT2RM4001454, NT2RM4001489, NT2RM4001605, NT2RM4001611, NT2RM4001666, NT2RM4001710, NT2RM4001714, NT2RM4001715, NT2RM4001731, NT2RM4001741, NT2RM4001746, NT2RM4001856, NT2RM4001938, NT2RM4001940, NT2RM4001984,

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	NT2RP4001041,	NT2RP4001100,	NT2RP4001117,	NT2RP4001174,	NT2RP4001235,	NT2RP4001274,
	NT2RP4001276,	NT2RP4001315,	NT2RP4001373,	NT2RP4001389,	NT2RP4001433,	NT2RP4001502,
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[0095] These clones are associated with ultraviolet radiation damage.
      A survey of genes of which expression levels are varied in response to the stimulation for inducing cell dif-
ferentiation (stimulation using retinoic acid (RA) or using RA/inhibitor (inhibitor for cell division)) in culture cells of neural
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HEMBA1000005.

strain, NT2, revealed the following clones with elevated expression levels in the presence of RA:

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                                    PLACE2000450,
                                                    PLACE2000465,
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                                                                                    PLACE3000401,
     PLACE3000402,
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                                    PLACE3000455,
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     THYRO1000034.
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                                    THYRO1000092.
                                                   THYRO1000111,
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                    THYRO1000387,1
                                    THYRO1000394,
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                    THYRO1000793,
                                    THYRO1000796,
                                                   THYRO1000805,
                                                                   THYRO1000815,
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     THYRO1000916,
                    THYRO1000934,
                                    THYRO1000974,
                                                   THYRO1000975,
                                                                   THYRO1001031.
                                                                                   THYRO1001062,
     THYRO1001093,
                    THYRO1001133,
                                    THYRO1001173,
                                                    THYRO1001177.
                                                                   THYRO1001189,
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                                    THYRO1001290.
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     THYRO1001617.
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                                    THYRO1001706,
                                                    THYRO1001772,
                                                                   THYRO1001828.
                                                                                   THYRO1001854.
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                                    Y79AA1000355,
                                                    Y79AA1000410.
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                                    Y79AA1000850.
                                                    Y79AA1000962.
                                                                    Y79AA1000976.
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     Y79AA1001711,
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                                                    Y79AA1002258,
                                                                    Y79AA1002361,
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     Y79AA1002472, Y79AA1002482
     [0097] On the other hand, clones of which expression levels decreased by RA are as follows:
     HEMBA1000946.
                    HEMBA1003569,
                                    HEMBA1005570,
                                                   HEMBB1000915,
                                                                    NT2RM1000666,
                                                                                   NT2RM2000092,
     NT2RM2000594.
                    NT2RM2001256,
                                    NT2RM4001754,
                                                    NT2RM4001905,
                                                                    NT2RP2001675,
                                                                                    NT2RP2002047,
     NT2RP2005491,
                    NT2RP3000980,
                                    NT2RP3002081,
                                                    NT2RP3004594,
                                                                    NT2RP4001950,
                                                                                    NT2RP4002408,
     OVARC1000431.
                    OVARC1001942,
                                    OVARC1001943,
                                                    PLACE1003190,
                                                                    PLACE1004868,
                                                                                    PLACE1005923,
     PLACE1007257, PLACE1010624, Y79AA1000346
     [0098] Clones of which expression levels increase by RA/inhibitor are as follows:
     HEMBA1000046, HEMBA1000307, HEMBA1000434,
                                                   HEMBA1000504,
                                                                   HEMBA1000588.
                                                                                   HEMBA1000682,
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     HEMBA1000726,
                    HEMBA1000943,
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                                                    HEMBA1001094,
                                                                   HEMBA1001122,
                                                                                   HEMBA1001323,
     HEMBA1001361,
                    HEMBA1001455,
                                    HEMBA1001709.
                                                   HEMBA1001746,
                                                                   HEMBA1001869,
                                                                                   HEMBA1002084,
     HEMBA1002583,
                    HEMBA1002628.
                                    HEMBA1002801,
                                                   HEMBA1002937,
                                                                   HEMBA1003096.
                                                                                   HEMBA1003142.
     HEMBA1003229,
                    HEMBA1003276.
                                    HEMBA1003309,
                                                    HEMBA1003463.
                                                                   HEMBA1003597.
                                                                                   HEMBA1003617.
     HEMBA1003725,
                    HEMBA1003803,
                                    HEMBA1003879,
                                                    HEMBA1003989.
                                                                   HEMBA1004000.
                                                                                   HEMBA1004015,
50
     HEMBA1004024,
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                    HEMBA1004049.
                                    HEMBA1004056.
                                                                   HEMBA1004248.
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     HEMBA1004554.
                    HEMBA1004666.
                                    HEMBA1004725.
                                                                   HEMBA1004803,
                                                    HEMBA1004770.
                                                                                   HEMBA1004923,
     HEMBA1004934,
                    HEMBA1004954,
                                    HEMBA1005039,
                                                   HEMBA1005075,
                                                                   HEMBA1005113,
                                                                                   HEMBA1005219.
                                    HEMBA1005304,
     HEMBA1005232,
                    HEMBA1005251,
                                                   HEMBA1005367,
                                                                   HEMBA1005372,
                                                                                   HEMBA1005403,
     HEMBA1005410,
                    HEMBA1005411,
                                    HEMBA1005548,
                                                   HEMBA1005581,
                                                                   HEMBA1005631,
                                                                                   HEMBA1005666,
     HEMBA1005755,
                    HEMBA1005780,
                                    HEMBA1006067,
                                                   HEMBA1006130,
                                                                   HEMBA1006364.
                                                                                   HEMBA1006485.
     HEMBA1006559.
                    HEMBA1006579,
                                    HEMBA1006754,
                                                   HEMBB1000059,
                                                                   HEMBB1000575,
                                                                                   HEMBB1000709,
     HEMBB1000822,
                    HEMBB1000848,
                                    HEMBB1000852,
                                                    HEMBB1000913,
                                                                   HEMBB1000985,
                                                                                   HEMBB1001117,
     HEMBB1001210,
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                                                   HEMBB1001443,
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HEMBB1002266, HEMBB1002371, HEMBB1002502,
     HEMBB1002049.
                    HEMBB1002254.
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     HEMBB1002617.
                    HEMBB1002692, HEMBB1002697, MAMMA1000241, MAMMA1000424,
                                                                                   MAMMA1000616,
     MAMMA1000731, MAMMA1000824, MAMMA1000908, MAMMA1000956, MAMMA1001038,
                                                                                   MAMMA1001091.
     MAMMA1001243, MAMMA1001815, MAMMA1001820, MAMMA1002267, MAMMA1002769,
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                                     PLACE1011858,
                                                     PLACE1011923,
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                                                                                    PLACE2000039.
     PLACE2000216,
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     PLACE3000121,
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     PLACE4000522, PLACE4000558,
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                                                    THYRO1000596,
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                                     Y79AA1001402,
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     Y79AA1001827, Y79AA1001875, Y79AA1002027, Y79AA1002211, Y79AA1002234, Y79AA1002258
     [0099] On the other hand, clones of which expression levels decrease by RA/inhibitor are as follows:
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                    HEMBA1000501, HEMBA1000946,
     HEMBA1000012,
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                                    HEMBB1001665,
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                                                                     NT2RP1001073.
                                                                                    NT2RP2000098,
     NT2RP2000965,
                     NT2RP2001397,
                                     NT2RP2002047,
                                                     NT2RP2004226.
                                                                     NT2RP2004396,
                                                                                    NT2RP2004655.
     NT2RP2005126,
                     NT2RP2005464,
                                     NT2RP2005712,
                                                     NT2RP2005859,
                                                                     NT2RP2005890.
                                                                                    NT2RP3000980,
     NT2RP3001383,
                     NT2RP3001621,
                                     NT2RP3002081,
                                                     NT2RP3002181,
                                                                     NT2RP3002244,
                                                                                    NT2RP3002590,
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     NT2RP3003059,
                     NT2RP3004258,
                                     NT2RP3004378,
                                                     NT2RP3004527,
                                                                    NT2RP3004594,
                                                                                    NT2RP4001760,
     NT2RP4001950,
                     NT2RP4002047,
                                    NT2RP4002408,
                                                    NT2RP5003459,
                                                                    OVARC1000004.
                                                                                    OVARC1000035,
     OVARC1000431,
                     OVARC1001051,
                                     OVARC1001129,
                                                    OVARC1001176,
                                                                    OVARC1001261,
                                                                                    OVARC1001342,
     OVARC1001942.
                     OVARC1001943,
                                     PLACE1002171,
                                                     PLACE1002465,
                                                                     PLACE1003190,
                                                                                    PLACE1003375,
     PLACE1004128,
                     PLACE1005026,
                                     PLACE1005876,
                                                     PLACE1005923,
                                                                     PLACE1007257,
                                                                                    PLACE1007375,
     PLACE1007507,
                     PLACE1008941,
                                     PLACE1010624,
                                                    PLACE1011090,
                                                                    PLACE1011219,
                                                                                    THYRO1000270,
     Y79AA1000346, Y79AA1001541
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[0100] These clones are also associated with neural differentiation and, therefore, are candidates for genes associated with neurological diseases.

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[0101] For example, if the protein encoded by the cDNA of the present invention is a regulatory factor of cellular conditions such as growth and differentiation, it can be used for developing medicines as follows. The protein or antibody provided by the invention is injected into a certain kind of cells by microinjection. Then, using the cells, it is possible to screen low molecular weight compounds by measuring the change in the cellular conditions, or the activation or inhibition of a particular gene. The screening can be performed as follows. First, the protein is expressed and purified as recombinant. The purified protein is microinjected into cells such as various cell lines, or primary culture cells, and the cellular change such as growth and differentiation can be examined. Alternatively, the induction of genes whose expression is known to be associated with a particular change of cellular conditions may be detected by the amount of mRNA or protein. Or, the amount of intracellular molecules (low molecular weight compounds, etc.) that is changed by the function of the gene product (protein) which is known to be associated with a particular change of cellular conditions may be detected. The compounds to be screened (both low and high molecular compounds are acceptable) can be added to the culture media and assessed for their activity by measuring the change of the cellular conditions. Instead of microinjection, cell lines introduced with the gene obtained in the invention can be used for the screening. If the gene product is turn out to be associated with a particular change in the cellular conditions, the change of the product can be used as a measurement for screening. Once a compound is screened out which can activate or inhibit

the function of the protein of the invention, it can be applied for developing medicines.

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[0102] If the protein encoded by the cDNA of the present invention is a secretory protein, membrane protein, or protein associated with signal transduction, glycoprotein, transcription, or diseases, it can be used in functional assays for developing medicines.

[0103] In case of a membrane protein, it is most likely to be a protein that functions as a receptor or ligand on the cell surface. Therefore, it is possible to reveal a new relationship between a ligand and receptor by screening the membrane protein of the invention based on the binding activity with the known ligand or receptor. Screening can be performed according to the known methods.

[0104] For example, a ligand against the protein of the invention can be screened in the following manner. Namely, a ligand that binds to a specific protein can be screened by a method comprising the steps of: (a) contacting a test sample with the protein of the invention or a partial peptide thereof, or cells expressing these, and (b) selecting a test sample that binds to said protein, said partial peptide, or said cells.

[0105] On the other hand, for example, screening using cells expressing the protein of the present invention that is a receptor protein can also be performed as follows. It is possible to screen receptors that is capable of binding to a specific protein by using procedures (a) attaching the sample cells to the protein of the invention or its partial peptide, and (b) selecting cells that can bind to the said protein or its partial peptide.

[0106] In a following screening as an example, first the protein of the invention is expressed, and the recombinant protein is purified. Next, the purified protein is labeled, binding assay is performed using a various cell lines or primary cultured cells, and cells that are expressing a receptor are selected (Growth and differentiation factors and their receptors, Shin-Seikagaku Jikken Kouza Vol.7 (1991) Honjyo, Arai, Taniguchi, and Muramatsu edit, p203-236, Tokyo-Kagaku-Doujin). A protein of the invention can be labeled with RI such as ¹²⁵I, and enzyme (alkaline phosphatase etc.). Alternatively, a protein of the invention may be used without labeling and then detected by using a labeled antibody against the protein. The cells that are selected by the above screening methods, which express a receptor of the protein of the invention, can be used for the further screening of an agonists or antagonists of the said receptor.

[0107] Once the ligand binding to the protein of the invention, the receptor of the protein of the invention or the cells expressing the receptor are obtained by screening, it is possible to screen a compound that binds to the ligand and receptor. Also it is possible to screen a compound that can inhibit both bindings (agonists or antagonists of the receptor, for example) by utilizing the binding activities.

[0108] When the protein of the invention is a receptor, the screening method comprises the steps of (a) contacting the protein of the invention or cells expressing the protein of the invention with the ligand, in the presence of a test sample, (b) detecting the binding activity between said protein or cells expressing said protein and the ligand, and (c) selecting a compound that reduces said binding activity when compared to the activity in the absence of the test sample. Furthermore, when the protein of the invention is a ligand, the screening method comprises the steps of (a) contacting the protein of the invention with its receptor or cells expressing the receptor in the presence of samples, (b) detecting the binding activity between the protein and its receptor or the cells expressing the receptor, and (c) selecting a compound that can potentially reduce the binding activity compared to the activity in the absence of the sample.

[0109] Samples to screen include cell extracts, expressed products from a gene library, synthesized low molecular compound, synthesized peptide, and natural compounds, for example, but are not construed to be listed here. A compound that is isolated by the above screening using a binding activity of the protein of the invention can also be used as a sample.

[0110] A compound isolated by the screening may be a candidate to be an agonist or an antagonist of the receptor of the protein. By utilizing an assay that monitors a change in the intracellular signaling such as phosphorylation which results from reduction of the binding between the protein and its receptor, it is possible to identify whether the obtained compound is an agonist or antagonist of the receptor. Also, the compound may be a candidate of a molecule that can inhibit the interaction between the protein and its associated proteins (including a receptor) in vivo. Such compounds can be used for developing drugs for precaution or cures of a disease with which the protein is associated.

[0111] Secretory proteins may regulate cellular conditions such as growth and differentiation. It is possible to find out a novel factor that regulates cellular conditions by adding the secretory protein of the invention to a certain kind of cell, and performing a screening by utilizing the cellular changes in growth or differentiation, or activation of a particular gene.

[0112] The screening can be performed, for example, as follows. First, the protein of the invention is expressed and purified in a recombinant form. Then, the purified protein is added to a various kind of cell lines or primary cultured cells, and the change in the cell growth and differentiation is monitored. The induction of a particular gene that is known to be involved in a certain cellular change is detected by the amounts of mRNA and protein. Alternatively, the amount of an intracellular molecule (low-molecular-weight compounds, etc.) that is changed by the function of a gene product (protein) that is known to function in a certain cellular change is used for the detection.

[0113] Once the screening reveals that the protein of the invention can regulate cellular conditions or the functions, it is possible to apply the protein as a pharmaceutical and diagnostic medicine for associated diseases by itself or by

altering a part of it into an appropriate composition.

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[0114] As is above described for membrane proteins, the secretory protein provided by the invention may be used to explore a novel ligand-receptor interaction using a screening based on the binding activity to a known ligand or receptor. A similar method can be used to identify an agonist or antagonist. The resulting compounds obtained by the methods can be a candidate of a compound that can inhibit the interaction between the protein of the invention and an interacting molecule (including a receptor). The compounds may be able to use as a preventive, therapeutic, and diagnostic medicine for the diseases, in which the protein may play a certain role.

[0115] Proteins associated with signal transduction or transcription may be a factor that affects a certain protein or gene in response to intracellular/extracellular stimuli. It is possible to find out a novel factor that can affect a protein or gene by expressing the protein provided by the invention in a certain types of cells, and performing a screening utilizing the activation of a certain intracellular protein or gene.

[0116] The screening may be performed as follows. First, a transformed cell line expressing the protein is obtained. Then, the transformed cell line and the untransformed original cell line are compared for the changes in the expression of a certain gene by detecting the amount of its mRNA or protein. Alternatively, the amount of an intracellular molecule (low molecular weight compounds) that is changed by the function of a certain gene product (protein) may be used for the detection. Furthermore, the change of the expression of a certain gene can be detected by introducing a fusion gene that comprises a regulatory region of the gene and a marker gene (luciferase, beta-galactosidase, etc.) into a cell, expressing the protein provided by the invention into the cell, and estimating the activity of a marker gene product (protein).

[0117] If the protein or gene of the invention is associated with diseases, it is possible to screen a gene or compound that can regulate its expression and/or activity either directly or indirectly by utilizing the protein of the present invention.

[0118] For example, the protein of the invention is expressed and purified as a recombinant protein. Then, the protein or gene that interacts with the protein of the invention is purified, and screened based on the binding. Alternatively, the screening can be performed by adding with a compound of a candidate of the inhibitor added in advance and monitoring the change of binding activity. In another method, a transcription regulatory region locating in the 5'-upstream of the gene encoding the protein of the invention that is capable of regulating the expression of other genes is obtained, and fused with a marker gene. The fusion is introduced into a cell, and the cell is added with compounds to explore a regulatory factor of the expression of the said gene.

[0119] The compound obtained by the screening can be used for developing pharmaceutical and diagnostic medicines for the diseases with which the protein of the present invention is associated. Similarly, if the regulatory factor obtained in the screening is turn out to be a protein, compounds that can newly affect the expression or activity of the protein may be used as a medicine for the diseases with which the protein of the invention is associated.

[0120] If the protein of the invention has an enzymatic activity, regardless as to whether it is a secretory protein, membrane protein, or proteins associated with signal transduction, glycoprotein, transcription, or diseases, a screening may be performed by adding a compound to the protein of the invention and monitoring the change of the compound. The enzymatic activity may also be utilized to screen a compound that can inhibit the activity of the protein.

[0121] In a screening given as an example, the protein of the invention is expressed and the recombinant protein is purified. Then, compounds are contacted with the purified protein, and the amount of the compound and the reaction products is examined. Alternatively, compounds that are candidates of an inhibitor are pretreated, then a compound (substrate) that can react with the purified protein is added, and the amount of the substrate and the reaction products is examined.

[0122] The compounds obtained in the screening may be used as a medicine for diseases with which the protein of the invention is associated. Also they can be applied for tests that examine whether the protein of the invention functions normally *in vivo*.

[0123] Whether the secretory protein, membrane protein, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein of the present invention is a novel protein associated with diseases or not is determined in another method than described above, by obtaining a specific antibody against the protein of the invention, and examining the relationship between the expression or activity of the protein and a certain disease. In an alternative way, it may be analyzed referred to the methods in "Molecular Diagnosis of Genetic Diseases" (Elles R. edit, (1996) in the series of "Method in Molecular Biology" (Humana Press).

[0124] Proteins associated with diseases are targets of screening as mentioned, and thus are very useful in developing drugs which regulate their expression and activity. Also, the proteins are useful in the medicinal industry as a diagnostic marker of the associated disease or a target of gene therapy.

[0125] Compounds isolated as mentioned above can be administered patients as it is, or after formulated into a pharmaceutical composition according to the known methods. For example, a pharmaceutically acceptable carrier or vehicle, specifically sterilized water, saline, plant oil, emulsifier, or suspending agent can be mixed with the compounds appropriately. The pharmaceutical compositions can be administered to patients by a method known to those skilled in the art, such as intraarterial, intravenous, or subcutaneous injections. The dosage may vary depending on the weight

or age of a patient, or the method of administration, but those skilled in the art can choose an appropriate dosage properly. If the compound is encoded by DNA, the DNA can be cloned into a vector for gene therapy, and used for gene therapy. The dosage of the DNA and the method of its administration may vary depending on the weight or age of a patient, or the symptoms, but those skilled in the art can choose properly.

[0126] The present invention further relates to databases comprising at least a sequence of polynucleotide and/or protein, or a medium recorded in such databases, selected from the sequence data of the nucleotide and/or the amino acids indicated in Table 350 and Table 351.

The term "database" means a set of accumulated information as machine-searchable and readable information of nucleotide sequence. The databases of the present invention comprise at least one of the novel nucleotide sequences of polynucleotides provided by the present invention. The databases of the present invention can consist of only the sequence data of the novel polynucleotides provided by the present invention or can comprise other information on nucleotide sequences of known full-length cDNAs or ESTs. The databases of the present invention can be comprised of not only the information on the nucleotide sequences but also the information on the gene functions revealed by the present invention. Additional information such as names of DNA clones carrying the full-length cDNAs can be recorded or linked together with the sequence data in the databases.

[0127] The database of the present invention is useful for gaining complete gene sequence information from partial sequence information of a gene of interest. The database of the present invention comprises nucleotide sequence information of full-length cDNAs. Consequently, by comparing the information in this database with the nucleotide sequence of a partial gene fragment yielded by differential display method or subtraction method, the information on the full-length nucleotide sequence of interest can be gained from the sequence of the partial fragment as a starting clue.

[0128] The sequence information of the full-length cDNAs constituting the database of the present invention contains not only the information on the complete sequences but also extra information on expression frequency of the genes as well as homology of the genes to known genes and known proteins. Thus the extra information facilitates rapid functional analyses of partial gene fragments. Further, the information on human genes is accumulated in the database of the present invention, and therefore, the database is useful for isolating a human homologue of a gene originating from other species. The human homologue can be isolated based on the nucleotide sequence of the gene from the original species.

[0129] At present, information on a wide variety of gene fragments can be obtained by differential display method and subtraction method. In general, these gene fragments are utilized as tools for isolating the full-length sequences thereof. When the gene fragment corresponds to an already-known gene, the full-length sequence is easily obtained by comparing the partial sequence with the information in known databases. However, when there exists no information corresponding to the partial sequence of interest in the known databases, cDNA cloning should be carried out for the full-length cDNA. It is often difficult to obtain the full-length nucleotide sequence using the partial sequence information as an initial clue. If the full-length of the gene is not available, the amino acid sequence of the protein encoded by the gene remains unidentified. Thus the database of the present invention can contribute to the identification of full-length cDNAs corresponding to gene fragments, which cannot be revealed by using databases of known genes.

[0130] The present invention has provided 5602 novel full-length cDNA clones, and primers for synthesizing the cDNA. As has not yet proceeded the isolation of full-length cDNA within the human, the invention has great significance. The full-length cDNA clones contain the translation initiation site, and thus provide a useful information for analysis of protein functions.

[0131] The cDNA clones are assumed to encode proteins such as secretory proteins, membrane proteins, signal transduction-associated protein, glycoprotein-associated protein, or transcription-associated protein, etc., which have important functions in vivo, and also predicted to be associated with many diseases. The genes and proteins associated with diseases are useful for developing a diagnostic marker or medicines for regulation of their expression and activity, or as a target of gene therapy.

[0132] The invention is illustrated more specifically with reference to the following examples, but is not to be construed as being limited thereto.

EXAMPLE 1

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Construction of a cDNA library by the oligo-capping method.

[0133] The NT-2 neuron progenitor cells (Stratagene), a teratocarcinoma cell line from human embryo testis, which can differentiate into neurons by the treatment with retinoic acid were used.

The NT-2 cells were cultured according to the manufacturer's instructions as follows.

- (1) NT-2 cells were cultured without induction by retinoic acid treatment (NT2RM1, NT2RM2, NT2RM4).
- (2) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 48 hours (NT2RP1).

(3) After cultured, NT-2 cells were induced by adding retinoic acid, and then were cultured for 2 weeks (NT2RP2, NT2RP3, NT2RP4, NT2RP5).

[0134] Also, the human neuroblastoma cell line SK-N-MC (ATCC HTB-10) (SKNMC1), and human retinoblastoma cell line Y79 (ATCC HTB-18) (Y79AA1) were cultured according to the culture conditions described in the ATCC catalogue (http://www.atcc.org/). The cells were harvested separately, and mRNA was extracted from each cell by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)+RNA was purified from the mRNA using oligo-dT cellulose. [0135] Similarly, human placenta (PLACE1, PLACE2, PLACE3, PLACE4), human ovary cancer tissue (OVARC1), tissues from human embryo at 10 weeks, which is enriched with head (HEMBA1), or body (HEMBB1), human mammary gland (MAMMA1), human thyroid gland (THYRO1), and primary cultured cells of human blood vessel endothelium (VESEN1) were used to extract mRNA by the method described in the literature (Molecular Cloning 2nd edition. (1989) Sambrook J., Fritsch, E.F., and Maniatis T., Cold Spring Harbor Laboratory Press). Furthermore, poly(A)+RNA was purified from the mRNA using oligo-dT cellulose.

[0136] Each poly(A)+RNA was used to construct a cDNA library by the oligo-capping method (Maruyama M. and Sugano S. (1994) Gene, 138: 171-174). Using the Oligo-cap linker (SEQ ID NO: 10464) and the Oligo-dT primer (SEQ ID NO: 10465), bacterial alkaline phosphatase (BAP) treatment, tobacco acid phosphatase (TAP) treatment, RNA ligation, the first strand cDNA synthesis, and removal of RNA were performed as described in the reference (Suzuki and Kanno (1996) Protein Nucleic acid and Enzyme, 41: 197-201; Suzuki Y. et al. (1997) Gene, 200: 149-156), Next, 5'- and 3'-PCR primers (SEQ ID NO: 10466, and 10467, respectively) were used for performing PCR to convert the cDNA into double stranded cDNA, which was then digested with Sfil. Then, the DrallI-cleaved pUC19FL3 vector (Figure 1; for NT2RM1, and NT2RP1), or the DrallI-cleaved pME18SFL3 (Figure 1) (GenBank AB009864, expression vector; for NT2RM2, NT2RM4, NT2RP2, NT2RP3, NT2RP4, NT2RP5, SKNMC1, Y79AA1, PLACE1, PLACE2, PLACE3, PLACE4, OVARC1, HEMBA1, HEMBB1, MAMMA1, THYRO1, and VESEN1) was used for cloning the cDNA in a unidirectional manner, and cDNA libraries were obtained. The nucleotide sequence of the 5'- and 3'- ends of the cDNA clones was analyzed with a DNA sequencer (ABI PRISM 377, PE Biosystems) after sequencing reactions were performed with the DNA sequencing reagents (Dye Terminator Cycle Sequencing FS Ready Reaction Kit, dRhodamine Terminator Cycle Sequencing FS Ready Reaction Kit, or BigDye Terminator Cycle Sequencing FS Ready Reaction Kit, PE Biosystems), according to the instructions. The data were compiled into a database.

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[0137] The full-length-enriched cDNA libraries except those for NT2RM1 and NT2RP1 were constructed using eukaryotic expression vector pME18SFL3. The vector contains SRα promoter and SV40 small t intron in the upstream of the cloning site, and SV40 polyA added signal sequence site in the downstream. As the cloning site of pME18SFL3 has asymmetrical DrallI sites, and the ends of cDNA fragments contain Sfil sites complementary to the DrallI sites. the cloned cDNA fragments can be inserted into the downstream of the SR α promoter unidirectionally. Therefore, clones containing full-length cDNA can be expressed transiently by introducing the obtained plasmid directly into COS cells. Thus, the clones can be analyzed very easily in terms of the proteins that are the gene products of the clones, or in terms of the biological activities of the proteins.

[0138] Herein, the cDNA libraries and the name of each clone are related as shown in Table 3. Therein, "xxxxxx" represents the clone number of six digits. Thus, the sequences are named by the library name, the clone number plus F- for the 5'-end, or R- for the 3'-end.

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Table 3

5	library: clone	5'-end sequence	3'-end sequence
10	NT2RM1: NT2RM1xxxxxx NT2RP1:	F-NT2RM1xxxxxx	
	NT2RP1xxxxxx NT2RM2:	F-NT2RP1xxxxxx	
15	NT2RM2xxxxxx NT2RM4:		
	NT2RM4xxxxxx	F-NT2RM4xxxxxx	R-NT2RM4xxxxxx
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	NT2RP2:		
	NT2RP2xxxxxx	F-NT2RP2xxxxxx	R-NT2RP2xxxxxx
5	NT2RP3:		
	NT2RP3xxxxxx	F-NT2RP3xxxxxx	R-NT2RP3xxxxxx
	NT2RP4:		
	NT2RP4xxxxxx	F-NT2RP4xxxxxx	R-NT2RP4xxxxxx
10	NT2RP5:		
	NT2RP5xxxxxx	F-NT2RP5xxxxxx	R-NT2RP5xxxxxx
	SKNMC1:		
	SKNMC1xxxxxx	F-SKNMC1xxxxxx	R-SKNMC1xxxxxx
15	Y79AA1:		
	Y79AA1xxxxxx	F-Y79AA1xxxxxx	R-Y79AA1xxxxxx
	PLACE1:		
20	PLACE1xxxxxx	F-PLACE1xxxxxx	R-PLACE1xxxxxx
	PLACE2:		
	PLACE2xxxxxx	F-PLACE2xxxxxx	R-PLACE2xxxxxx
	PLACE3:		
25	PLACE3xxxxxx	F-PLACE3xxxxxx	R-PLACE3xxxxxx
	PLACE4:		•
	PLACE4xxxxxx	F-PLACE4xxxxxx	R-PLACE4xxxxxx
	OVARC1:		•
30	OVARC1xxxxxx	F-0VARC1xxxxxx	R-OVARC1xxxxxx
	HEMBA1:		
	HEMBA1xxxxxx	F-HEMBA1xxxxxx	R-HEMBA1xxxxxx
35	HEMBB1:		
	HEMBB1xxxxxx	F-HEMBB1xxxxxx	R-HEMBB1xxxxxx
	MAMMA1:		O 144 104 A 4
	MAMMA1xxxxxx	F-MAMMA1xxxxxx	R-MAMMA1xxxxx
40	THYRO1:	E TUVD01	D TUVDA1
	THYR01xxxxxx	F-THYR01xxxxxx	R-THYR01xxxxxx
	VESEN1: VESEN1xxxxxx	F-VESEN1xxxxxx	R-VESEN1xxxxxx
	AESENIXXXXXX	L-AE9ENIXXXXXX	r_AE9EN (XXXXXX
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EXAMPLE 2

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Estimation of the fullness ratio at the 5'-ends of the clones contained in the cDNA libraries constructed by the oligocapping method.

[0139] The fullness ratio at the 5'-end sequence of the 59,823 clones in the human cDNA libraries constructed by the oligo-capping method was determined as follows. Of all the clones whose 5'-end sequences were found in those of known human mRNA in the public database, a clone was judged to be "full-length", if it had a longer 5'-end sequence than that of the known human mRNA, or, even though the 5'-end sequence was shorter, if it contained the translation initiation codon. A clone which did not contain the translation initiation codon was judged to be "not-full-length". The fullness ratio ((the number of full-length clones)/(the number of full-length and not-full-length clones)) at the 5'-end of the cDNA clones from each library was determined by comparing with the known human mRNA. As a result, the fullness

ratio of the 5'-ends was 63.5%. The result indicates that the fullness ratio at the 5'-end sequence was extremely high.

EXAMPLE 3

5 Assessment of the fullness ratio of the 5'-end of the cDNA by the ATGpr and the ESTiMateFL.

[0140] The ATGpr, developed by Salamov A.A., Nishikawa T., and Swindells M.B. in the Helix Research Institute, is a program for prediction of the translation initiation codon based on the characteristics of the sequences in the vicinity of the ATG codon. The results are shown with expectations (also described as ATGpr1 below) that an ATG is a true initiation codon (0.05-0.94) (can be described as ATGpr1). When the program was applied to the 5'-end sequences of the clones from the cDNA library that was obtained by the oligo-capping method and that had 65% fullness ratio, the sensitivity and specificity of estimation of a full-length clone (clone containing the N-terminal end of ORF) were improved to 82-83% by selecting only clones having the ATGpr1 score 0.6 or higher.

[0141] Furthermore, the program was used to assess the fullness of 18,959 clones in the human libraries obtained here, which have 5'-ends matched to a known human mRNA.

[0142] Briefly, the maximal ATGpr1 score of the clones was determined, and then their 5'-end sequence was compared with the known human mRNA to estimate whether the clone is full-length or not. The result is shown in Table 4. [0143] Based on the knowledge that known mRNAs, in general, are highly expressed in the cell, similar estimation was performed with genes having a low number in the EST hit, which represent relatively low abundant mRNAs, and the result is shown in Table 5.

[0144] In the table, the number of full-length clones indicate that of clones containing the N-terminal end of ORF, and so does the number of not-full-length clones that of clones without the N-terminal end of ORF. The fullness ratio represents (the number of full-length clones)/(the number of full-length clones).

Table 4

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having a matched 5'-end with that of a known mRNA.

maximal ATGpr1 score number of (full-length number of full-length fullness ratio clones plus not-full-length clones clones) >=0.70 11,193 9.346 83.5% >=0.50 13,369 10,549 78.9% >=0.30 15,489 11,340 73.2% >=0.15 17,394 11,811 67.9% >=0.00 18,959 12,046 63.5%

Table 5

The maximal ATGpr1 score and the fullness ratio of the 5'-end sequences of the clones obtained from human cDNA libraries constructed by the oligo-capping method; clones having 5 EST hits or less among the clones having a matched 5'-end with that of a known mRNA.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	2,801	1,934	69.0%
>=0.50	3,683	2,393	65.0%
>=0.30	4,683	2,707	57.8%
>=0.15	5,559	2,890	52.0%
>=0.00	6,113	3,013	49.8%

[0145] Next, the ESTiMateFL was used for the estimation. The ESTiMateFL, developed by Nishikawa and Ota in the Helix Research Institute, is a method for the selection of a clone with high fullness ratio by comparing with the 5'-end or 3'-end sequences of ESTs in the public database.

[0146] By the method, a cDNA clone is judged to be most likely not to be full-length if there exist any ESTs which

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have longer 5'-end or 3'-end sequences than the clone. The method is systematized for high throughput analysis. A clone is judged to be full-length if the clone has a longer 5'-end sequence than ESTs in the public database. Even if a clone has a shorter 5'-end, the clone is judged to be full-length if the difference in length is within 50 bases, and otherwise judged not to be full-length, for convenience.

[0147] In case of the clones whose 5'-end sequence is matching with the known mRNA, 80% of the clones judged to be full-length by comparing with ESTs was also judged to be full-length by comparing with the known mRNA. Also, 80% of the clones judged to be not full-length by comparing with ESTs was also judged to be not full-length by comparing with the known mRNA.

[0148] The precision of the estimation by comparing with ESTs is improved with increasing number of ESTs to be compared. However, in case that a limited number of ESTs are available, the reliability becomes low. Thus, the method is effective in excluding clones with high probability of being not-full-length, from the cDNA clones that is synthesized by the oligo-capping method and that have the 5'-end sequences with about 60 % fullness ratio. In particular, the ESTiMateFL is efficiently used to estimate the fullness ratio at the 3'-end sequence of cDNA of a human unknown mRNA which has a significant number of EST deposits in the public database.

[0149] The 18,959 clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, were estimated by using the ATGpr and ESTiMateFL. Briefly, the 5'-end sequence of the respective clone was analyzed to obtain the maximal ATGpr1 score, and compared with the ORF of the known human mRNA that matches with it to determine whether the clone is full-length or not. Then, the 5'-end sequence of the respective clone was analyzed by the ESTiMateFL to judge whether the clone is full-length or not. Specifically, the 5'-end sequences of the 18,959 clones were compared with those of ESTs by the ESTiMateFL and the clones other than those that are not full-length were selected. Then, the selected clones were used to analyze the relationship between the ATGpr and the fullness ratio. The result was summarized in Table 6. Also, among the selected, the clones in which the number of the EST hit is not more than 5 were selected and analyzed. The result was summarized in Table 7, which represents the result of the analysis of mRNA with relatively low abundance.

[0150] Therein, the number of being full-length, the number of being not full-length, and the fullness ratio indicate the number of the clones that contain the N-terminus of the ORF, the number of the clones that do not contain the N-terminus of the ORF, and (the number of being full-length)/(the number of being full-length) plus (the number of being not full-length), respectively.

Table 6

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The maximal ATGpr1 score and the fullness ratio of the 5'-end sequence in the clones isolated from human cDNA libraries constructed by the oligo-capping method, which have the 5'-end sequence matched with a known human mRNA, and also other than those being not full-length according to the comparison with ESTs.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	9,068	8,349	92.1%
>=0.50	10,345	9,318	90.1%
>=0.30	11,425	9,964	87.2%
>=0.15	12,254	10,335	84.3%
>=0.00	12,785	10,484	82.0%

Table 7

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.70	1,959	1,510	77.1%
>=0.50	2,469	1,821	73.8%
>=0.30	2,975	2,046	68.8%
>=0.15	3,368	2,164	64.3%

Table 7 (continued)

Maximal ATGpr1 score and fullness ratio of the 5'-end sequence of the clones, which were isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA, in which the number of the EST hit is not more than 5.

maximal ATGpr1 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.00	3,661	2,226	60.8%

[0151] The 19,226 clones, isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to that of a known human mRNA were estimated by the ATGpr2, and the correlation between the score and the fullness ratio was estimated. Specifically, the maximal ATGpr2 score of the clones identical to a known human mRNA was determined, and then their fullness ratio was estimated by comparing the 5'-ends with ORF of known human mRNA. The result was shown in Table 8.

Table 8

Maximal ATGpr2 score and fullness ratio of the 5'-end sequence of the clones, which are isolated from the human cDNA libraries constructed by the oligo-capping method, whose 5'-end sequence is identical to a known human mRNA.

maximal ATGpr2 score	number of (full-length clones plus not-full-length clones)	number of full-length clones	fullness ratio
>=0.30	10,748	8,031	74.7%
>=0.15	16,383	11,226	68.5%
>=0.00	19,226	12,285	63.9%

30 [0152] According to the above results, it was found that, in case of using clones isolated from human cDNA libraries constructed by the oligo-capping method, the fullness ratio of the clones that have low score in the ATGpr can be improved by estimating their 5'-end sequence using the combination of the ATGpr and the ESTiMateFL. Therefore, the method was applied to select a cDNA clone with high fullness ratio.

35 EXAMPLE 4

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Clustering of the 5'-end and 3'-end sequences of cDNA clones.

[0153] The 5'-end and 3'-end sequences of cDNA clones were obtained, and clustered separately. The single pass data of the nucleotide sequence of the 5'-end and 3'-end was subject to the BLAST search between the sequence data of all the clones synthesized in example 1, and the clones considered to be originating from the same gene were put together into a group. If the 5'-end of a clone contains the consensus sequence of 300 bases or more with identity 95% or more, or the 3'-end contains the consensus sequence of 200 bases or more and having identity 90% or more, the clones were put in the same group.

⁴⁵ **[0154]** The groups of the 5'-end sequence and the 3'-end sequence were further clustered so as that the groups from the same clone can be in the same group (cluster).

EXAMPLE 5

50 Characterization of the cloned sequence.

[0155] The data of the 5'-end sequence of the cloned sequence was characterized by the following way:

- (1) examining whether it is identical to those of mRNA from human and other species (including authorized sequences) and human EST by the BLAST homology search of the GenBank,
- (2) examining whether it has longer 5'-end than those of human mRNA and human EST,
- (3) determining the scores in the ATGpr1 and ATGpr2 programs of all the initiation codons in the 5'-end sequence, and

- (4) determining the number of the human EST clone(s) that is judged to be identical by the BLAST homology search of the GenBank.
- [0156] The data of the 3'-end sequence of the cloned sequence was characterized by the above (1) and (4).
- [0157] These characterized data were used for the final selection of the clones.

EXAMPLE 6

Identity to the human mRNA and human EST, and comparison of the length of the 5'-end.

[0158] The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of mRNA from human or other species when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 94% or more. The 5'-end and 3'-end sequences of the cloned sequence was judged to be identical to those of human EST when the sequence to compare has the length of 200 bases or longer, and the obtained homology is 90% or more.

[0159] The sequence of the clone was judged to be full-length in comparison with human mRNA when the sequence has longer 5'-end, or it contains the translation initiation site. The sequence of the clone was judged to be full-length in comparison with human EST in the database when the sequence has longer 5'-end, or while it has shorter end, the difference in length between the two sequences is 50 bases or less. The other clones were judged to be not full-length.

EXAMPLE 7

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Prediction of the fullness ratio by the ATGpr.

25 [0160] The score in the ATGpr1 is the expectation to be full-length based on calculations, and the higher score reflects the higher probability to be full-length as shown in Example 3. The maximal ATGpr1 score and the maximal ATGpr2 score represent the score obtained with all the initiation codons contained in the 5'-end sequence of the cloned sequence, and were used for the characterization.

30 EXAMPLE 8

Prediction of the novelty using the number of the identical ESTs by the homology search.

[0161] For both the 5'-end and 3'-end sequences of the clones, the number of the identical ESTs was determined by the homology search on the GenBank. Human ESTs were judged to be identical when the EST has a sequence of 200 nucleotides or more with 90% or more matching with the 5'-end sequence. The number of the identical ESTs were used for characterization and as an index of novelty. The clone having not identical sequence at the 5'-end and 3'-end sequences to those of mRNA as well as those of ESTs is a gene encoding a novel protein. Similarly, a clone having either the 5'-end or the 3'-end sequences, which has low number of the identical ESTs, is judged to be a gene encoding a novel protein.

EXAMPLE 9

Characterization of clusters.

[0162] The clusters of the groups of the 5'-end and 3'-end sequences were characterized according to the following criteria.

- (1) Whether it is identical to the mRNA sequences from human or other species (including authorized sequences), or human ESTs by the BLAST search of the GenBank.
- A cluster containing at least one sequence of all the 5'-end and 3'-end sequences, which is identical to one of the mRNA sequences, was regarded to be the same cluster of the mRNA sequence.
- (2) Whether it has longer 5'-end than human mRNA sequence and human ESTs.
- When all the 5'-end sequences contained in a cluster are judged to be not full-length compared with the mRNA sequences and human ESTs, the cluster was regarded as being not full-length.
- (3) The scores in the ATGpr1 and ATGpr2 using all the initiation codons contained in the 5'-end sequences. The maximal ATGpr1 score among those of all the 5'-end sequences in a cluster was determined as the ATGpr1 score of the cluster. The ATGpr2 score of the cluster was also determined in the same way.

(4) The number of the identical human ESTs determined by the BLAST search of the GenBank.

[0163] The maximum number was determined in the numbers of ESTs identical to each of 5'-end sequences contained in a cluster. The number of the ESTs identical to the 5'-end sequences in the cluster was defined as the maximum number. The number of the ESTs identical to the 3'-end sequences in a cluster was determined in the same way.

EXAMPLE 10

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Methods for selection of the clusters by the characteristics.

[0164] Data obtained by the characterization described above was used to discard the clusters that are identical to any mRNA sequence from human and other species (including authorized sequences), or those clusters that are not full-length. From the rest of the clusters, the clusters that fulfill any of the following conditions were selected.

- (a) A cluster in which the number of the identical ESTs for the 5'-end sequence is 20 or less, and the ATGpr1 score exceeds 0.3.
- (b) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for both the 5'-end sequence and the 3'-end sequence is 5 or less, and multiple clones are contained.
- (c) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is 0, and the number of the identical ESTs for the 3'-end sequence is not less than 1.
- (d) A cluster having the ATGpr1 score not more than 0.3, in which the number of the identical ESTs for the 5'-end sequence is not less than 1 and not more than 5, and the number of the identical ESTs for the 3'-end sequence is 0.

[0165] The clusters selected by (a) contain at least one clone that is novel and having high fullness ratio. The clusters selected by (b), (c), and (d) contain at least one clone that is novel and having low fullness ratio, but is still full-length.

EXAMPLE 11

Methods for selection of clones from clusters.

[0166] In the clusters comprising a single clone, the clone was selected.

[0167] In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score higher than 0.3, a clone with the highest score was selected.

[0168] In the clusters comprising multiple clones, in which multiple clones have the ATGpr1 score not more than 0.3, a clone with the highest ATGpr2 score was selected, if the score was higher than 0.3.

[0169] In the clusters comprising multiple clones, in which the clones have the scores not more than 0.3 in both the ATGpr1 and the ATGpr2, a clone with the highest scores in both the ATGpr1 and ATGpr2 was selected.

[0170] In the clusters comprising multiple clones, in which the above selection by the ATGpr score was not applicable, selected was a clone having longer 5'-end by assembling the 5'-end sequence, 3'-end sequence, and human ESTs. For assembling, the Sequencher™ (Hitachi Soft Engineering) was used. When even the selection by assembling failed, all the clones were judged to be full-length.

[0171] As a result, 3690 clones were the clones that have the maximal ATGpr1 score higher than 0.3. On the other hand, 477 clones were the clones that have the maximal ATGpr1 score not more than 0.3, and the maximal ATGpr2 score higher than 0.3. The number of the clones having the highest scores in both the ATGpr1 and ATGpr2, while the scores were not more than 0.3, were 97. The number of the clones which were not selected by the ATGpr scores, but were selected by assembling the 5'-end sequence, 3'-end sequence, and human ESTs, were 117. The clones that have the score in both the ATGpr1 and ATGpr2 not more than 0.3, but were selected because the cluster comprises a single clone, were 1166. In the clones, at least either of the 5'-end or 3'-end sequence was not identical to any of human ESTs. Some clones were selected because the cluster comprises a single clone, or by assembling, in which there is no ATG codon (9 clones: HEMBA1001960, HEMBA106569, HEMBB1001454, NT2PR2002839, NT2RP2005325, NT2RP2006323, PLACE1004506, PLACE1005526, and THYRO1001177). The sequences that do not contain the ATG codon were considered to be corresponding to the 5'-UTR. Although the clones do not have the scores in the ATGpr1 and ATGpr2, the clones were yet judged to be full-length according to the fullness ratio, as shown in Table 4, 5, 6, 7, and 8. The above clones that were finally judged to be full-length were classified into 11 groups according to the following criteria.

Group (1): 1516 clones

Among the 3690 clones having the maximal ATGpr1 score higher than 0.3, the following 1516 clones were having

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high fullness ratio and a novel clone, in which at least either of the 5'-end or 3'-end sequence, or both of them were
not identical to any of human ESTs.
HEMBA1000046, HEMBA1000050, HEMBA1000129, HEMBA1000150, HEMBA1000158, HEMBA1000193
HEMBA1000201, HEMBA1000216, HEMBA1000227, HEMBA1000288, HEMBA1000290, HEMBA100030 3,
HEMBA1000304, HEMBA1000369, HEMBA1000392, HEMBA1000396, HEMBA1000488, HEMBA10005 05,
HEMBA1000508, HEMBA1000534, HEMBA1000542, HEMBA1000594, HEMBA1000637, HEMBA1000 657,
HEMBA1000752, HEMBA1000867, HEMBA1000869, HEMBA1000872, HEMBA1000910, HEMBA100 0918.
HEMBA1000919, HEMBA1000942, HEMBA1000968, HEMBA1000975, HEMBA1000986, HEMBA10 01022
HEMBA1001043, HEMBA1001052, HEMBA1001080, HEMBA1001085, HEMBA1001109, HEMBA1 001140.
HEMBA1001174, HEMBA1001235, HEMBA1001286, HEMBA1001302, HEMBA1001398, HEMBA 1001407,
HEMBA1001415, HEMBA1001446, HEMBA1001476, HEMBA1001497, HEMBA1001510, HEMB A1001533,
HEMBA1001570, HEMBA1001581, HEMBA1001635, HEMBA1001640, HEMBA1001647, HEM BA1001661,
HEMBA1001731, HEMBA1001744, HEMBA1001746, HEMBA1001800, HEMBA1001815, HE MBA1001822,
HEMBA1001866, HEMBA1001896, HEMBA1001910, HEMBA1001987,
HEMBA1002018, HEMBA1002035, HEMBA1002049, HEMBA1002092, HEMBA1002119, HEMBA1002125,
HEMBA1002161, HEMBA1002177, HEMBA1002189, HEMBA1002191, HEMBA1002199, HEMBA100222 9,
HEMBA1002237, HEMBA1002265, HEMBA1002363, HEMBA1002417, HEMBA1002419, HEMBA10024 30,
HEMBA1002439, HEMBA1002477, HEMBA1002503, HEMBA1002508, HEMBA1002515, HEMBA1002 547,
HEMBA1002688, HEMBA1002703, HEMBA1002746, HEMBA1002750, HEMBA1002850, HEMBA100 2973.
HEMBA1003021, HEMBA1003067, HEMBA1003077, HEMBA1003078, HEMBA1003079, HEMBA10 03117,
HEMBA1003129, HEMBA1003175, HEMBA1003199, HEMBA1003235, HEMBA1003250, HEMBA1 003257,
HEMBA1003291, HEMBA1003322, HEMBA1003327, HEMBA1003370, HEMBA1003380, HEMBA 1003395,
HEMBA1003402, HEMBA1003461, HEMBA1003480, HEMBA1003538, HEMBA1003545, HEMB A1003556,
HEMBA1003581, HEMBA1003621, HEMBA1003645, HEMBA1003667, HEMBA1003720, HEM BA1003760,
HEMBA1003799, HEMBA1003807, HEMBA1003827, HEMBA1003836, HEMBA1003866, HE MBA1003879,
HEMBA1003880, HEMBA1003985, HEMBA1003989,
HEMBA1004011, HEMBA1004048, HEMBA1004056, HEMBA1004074, HEMBA1004133, HEMBA1004146,
HEMBA1004150, HEMBA1004199, HEMBA1004200, HEMBA1004238, HEMBA1004246, HEMBA100427 5,
HEMBA1004286, HEMBA1004289, HEMBA1004321, HEMBA1004327, HEMBA1004335, HEMBA10043 41,
HEMBA1004372, HEMBA1004389, HEMBA1004479, HEMBA1004499, HEMBA1004507, HEMBA1004 542,
HEMBA1004554, HEMBA1004573, HEMBA1004596, HEMBA1004632, HEMBA1004697, HEMBA100 4705,
HEMBA1004709, HEMBA1004711, HEMBA1004736, HEMBA1004751, HEMBA1004752, HEMBA10 04753,
HEMBA1004756, HEMBA1004758, HEMBA1004768, HEMBA1004771, HEMBA1004795, HEMBA1 004806,
HEMBA1004850, HEMBA1004863, HEMBA1004889, HEMBA1004923, HEMBA1004929, HEMBA 1004930,
HEMBA1004944, HEMBA1004972, HEMBA1004980, HEMBA1005019, HEMBA1005035, HEMB A1005050,
HEMBA1005066, HEMBA1005075, HEMBA1005079, HEMBA1005083, HEMBA1005113, HEM BA1005133,
HEMBA1005149, HEMBA1005219, HEMBA1005331, HEMBA1005338, HEMBA1005367, HE MBA1005411,
HEMBA1005468, HEMBA1005469, HEMBA1005517, HEMBA1005518, HEMBA1005526, H EMBA1005548,
HEMBA1005576, HEMBA1005583, HEMBA1005595, HEMBA1005609, HEMBA1005685, HEMBA1005732,
HEMBA1005755, HEMBA1005813, HEMBA1005815, HEMBA1005834, HEMBA1005884, HEMBA1005963,
HEMBA1005991,
HEMBA1006031, HEMBA1006100, HEMBA1006121, HEMBA1006138, HEMBA1006173, HEMBA1006182,
HEMBA1006198, HEMBA1006252, HEMBA1006272, HEMBA1006278, HEMBA1006291, HEMBA100629 3,
HEMBA1006344, HEMBA1006349, HEMBA1006377, HEMBA1006381, HEMBA1006398, HEMBA10064 24,
HEMBA1006467, HEMBA1006474, HEMBA1006483, HEMBA1006492, HEMBA1006494, HEMBA1006 497,
HEMBA1006502, HEMBA1006530, HEMBA1006579, HEMBA1006583, HEMBA1006643, HEMBA100 6674,
HEMBA1006682, HEMBA1006709, HEMBA1006717, HEMBA1006737, HEMBA1006754, HEMBA10 06758,
HEMBA1006767, HEMBA1006795, HEMBA1006796, HEMBA1006807, HEMBA1006832, HEMBA1 006900,
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HEMBB1000008, HEMBB1000024, HEMBB1000025, HEMBB1000083, HEMBB1000103, HEMBB1000173,
HEMBB1000175, HEMBB1000198, HEMBB1000240, HEMBB1000244, HEMBB1000338, HEMBB100033 9,
HEMBB1000391, HEMBB1000438, HEMBB1000510, HEMBB1000550, HEMBB1000556, HEMBB10005 89,
HEMBB1000591, HEMBB1000593, HEMBB1000632, HEMBB1000671, HEMBB1000673, HEMBB1000 693,
HEMBB1000706, HEMBB1000725, HEMBB1000781, HEMBB1000810, HEMBB1000826, HEMBB100 0835,
HEMBB1000848, HEMBB1000852, HEMBB1000870, HEMBB1000887, HEMBB1000908, HEMBB10 00927,
HEMBB1000947, HEMBB1000973, HEMBB1000991, HEMBB1001011, HEMBB1001014, HEMBB1 001020,
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HEMBB1001024, HEMBB1001058, HEMBB1001096, HEMBB1001105, HEMBB1001126, HEMBB 1001169,
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HEMBB1001394, HEMBB1001410, HEMBB1001426, HEMBB1001449, HEMBB1001482, HEM BB1001531,
HEMBB1001562, HEMBB1001564, HEMBB1001585, HEMBB1001635, HEMBB1001653, HE MBB1001665,
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HEMBB1002044, HEMBB1002050, HEMBB1002068, HEMBB1002142, HEMBB1002152, HEMBB1002193
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MAMMA1000713, MAMMA1000731, MAMMA1000746, MAMMA1000760, MAMMA10 00776, MAMMA1000842,
MAMMA1000843, MAMMA1000856, MAMMA1000865, MAMMA1000875, MAMMA1 000897, MAMMA1000906,
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MAMMA1002384, MAMMA1002413, MAMMA1002427, MAMMA1002454, MAMMA1002524, MAMMA100 2554,
MAMMA1002590, MAMMA1002617, MAMMA1002655, MAMMA1002665, MAMMA1002671, MAMMA10 02673.
MAMMA1002685, MAMMA1002699, MAMMA1002711, MAMMA1002782, MAMMA1002807, MAMMA1 002842,
MAMMA1002843, MAMMA1002868, MAMMA1002881, MAMMA1002886, MAMMA1002895, MAMMA 1002938,
MAMMA1002964, MAMMA1003013, MAMMA1003015, MAMMA1003044, MAMMA1003049, MAMM A1003056,
MAMMA1003066, MAMMA1003099, MAMMA1003104, MAMMA1003127,
NT2RM1000018, NT2RM1000059, NT2RM1000118, NT2RM1000131, NT2RM1000186, NT2RM1000242,
NT2RM1000252, NT2RM1000260, NT2RM1000271, NT2RM1000314, NT2RM1000318, NT2RM100035 4,
NT2RM1000399, NT2RM1000421, NT2RM1000563, NT2RM1000623, NT2RM1000666, NT2RM10007 25,
NT2RM1000742, NT2RM1000772, NT2RM1000802, NT2RM1000811, NT2RM1000850, NT2RM1000 885,
NT2RM1000894, NT2RM1000978, NT2RM1001059, NT2RM1001066, NT2RM1001072, NT2RM100 1082,
NT2RM1001085, NT2RM1001092, NT2RM1001115,
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Group (2): 377 clones

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Among the 3690 clones, the following 377 clones were full-length, and a novel clone, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5. HEMBA1000156, HEMBA1000231, HEMBA1000244, HEMBA1000302, HEMBA1000523, HEMBA1000531, HEMBA1000555, HEMBA1000568, HEMBA1000971, HEMBA1001009, HEMBA1001133, HEMBA100113 7, HEMBA1001257, HEMBA1001281, HEMBA1001289, HEMBA1001310, HEMBA1001377, HEMBA10013 88, HEMBA1001413, HEMBA1001455, HEMBA1001579, HEMBA1001702, HEMBA1001781, HEMBA1001 804, HEMBA1001819. HEMBA1001864, HEMBA1001869, HEMBA1001912, HEMBA1001915, HEMBA100 2084, HEMBA1002102, HEMBA1002212, HEMBA1002495, HEMBA1002513, HEMBA1002863, HEMBA10 02935, HEMBA1002970, HEMBA1002971, HEMBA1002997, HEMBA1002999, HEMBA1003035, HEMBA1 003041, HEMBA1003433, HEMBA1003568, HEMBA1003680, HEMBA1003692, HEMBA1003804, HEMBA 1003893, HEMBA1003953, HEMBA1004097, HEMBA1004207, HEMBA1004502, HEMBA1004560, HEMB A1004638, HEMBA1004693, HEMBA1004973, HEMBA1004977, HEMBA1005223, HEMBA1005244, HEM BA1005314, HEMBA1005680, HEMBA1005737, HEMBA1005746, HEMBA1005822, HEMBA1005891, HE MBA1006445, HEMBA1006535, HEMBA1006566, HEMBA1006877, HEMBA1006936, HEMBA1007178, H EMBA1007251, HEMBA1007267, HEMBB1000030, HEMBB1000215, HEMBB1000226, HEMBB1000449, HEMBB1000789, HEMBB1000975, HEMBB1001153, HEMBB1001175, HEMBB1001234, HEMBB1001387, HEMBB1001905, HEMBB100190 6, HEMBB1002300, HEMBB1002383, HEMBB1002489, HEMBB1002607, HEMBB1002705, MAMMA1000257, MAMMA1000416, MAMMA1000831, MAMMA1000841, MAMMA1001021, MAMMA1001041, MAMMA1001105, MAMMA1001139, MAMMA1001198, MAMMA1001730, MAMMA1001837, MAMMA100219 8, MAMMA1002209, MAMMA1002219, MAMMA1002292, MAMMA1002297, MAMMA1002637, MAMMA10026 50, MAMMA1002796, MAMMA1002987, MAMMA1003135, MAMMA1003150,

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50 Group (3): 1797 clones

Among the 3690 clones, the following 1797 clones were full-length, and novel clones, in which the number of the identical human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).

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Group (4): 453 clones

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Among the 1857 clones having the maximal ATGpr1 score not more than 0.3 (including the 9 clones whose 5'-end sequence does not contain the ATG codon), the following 453 clones were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.

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    HEMBA1006949, HEMBA1007078, HEMBA1007129, HEMBA1007147, HEM BA1007206, HEMBA1007279,
    HEMBA1007327,
    HEMBB1000005, HEMBB1000055, HEMBB1000144, HEMBB1000258, HEMBB1000318, HEMBB1000335,
    HEMBB1000354, HEMBB1000374, HEMBB1000402, HEMBB1000404, HEMBB1000480, HEMBB100049 3,
    HEMBB1000554, HEMBB1000573, HEMBB1000649, HEMBB1000652, HEMBB1000709, HEMBB10007 49,
    HEMBB1000790, HEMBB1000827, HEMBB1000831, HEMBB1000893, HEMBB1001004, HEMBB1001 008.
    HEMBB1001047, HEMBB1001315, HEMBB1001317, HEMBB1001326, HEMBB1001367, HEMBB100 1424.
    HEMBB1001436, HEMBB1001458, HEMBB1001535, HEMBB1001565, HEMBB1001747, HEMBB10 01749,
    HEMBB1001797, HEMBB1001836, HEMBB1001863, HEMBB1001875, HEMBB1001911, HEMBB1 001922,
    HEMBB1001925, HEMBB1001944, HEMBB1001983, HEMBB1001996, HEMBB1001997, HEMBB 1002092,
    HEMBB1002247, HEMBB1002266, HEMBB1002387, HEMBB1002425, HEMBB1002458, HEMB B1002522,
    HEMBB1002534, HEMBB1002582, HEMBB1002596, HEMBB1002617, HEMBB1002702, MAMMA1000043,
    MAMMA1000092, MAMMA1000129, MAMMA1000198, MAMMA1000221, MAMMA1000307, MAMMA1000331,
    MAMMA1000360, MAMMA1000402, MAMMA1000414, MAMMA1000500, MAMMA100052 2, MAMMA1000576,
    MAMMA1000594, MAMMA1000597, MAMMA1000720, MAMMA1000775, MAMMA10007 78, MAMMA1000798,
    MAMMA1000862, MAMMA1000876, MAMMA1000931, MAMMA1000940, MAMMA1000 941, MAMMA1000975,
    MAMMA1001038, MAMMA1001186, MAMMA1001220, MAMMA1001256, MAMMA100 1274, MAMMA1001341,
    MAMMA1001397, MAMMA1001420, MAMMA1001547, MAMMA1001670, MAMMA10 01679, MAMMA1001711,
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MAMMA1001745, MAMMA1001760, MAMMA1001769, MAMMA1001815, MAMMA1 001907, MAMMA1002056,
    MAMMA1002078, MAMMA1002093, MAMMA1002125, MAMMA1002132, MAMMA 1002145, MAMMA1002250,
    MAMMA1002311, MAMMA1002411, MAMMA1002498, MAMMA1002571, MAMM A1002701, MAMMA1002727,
    MAMMA1002728, MAMMA1002746, MAMMA1002764, MAMMA1002765, MAM MA1002820, MAMMA1002830.
    MAMMA1002909, MAMMA1002941, MAMMA1002973, MAMMA1003004, MA MMA1003007, MAMMA1003039.
    MAMMA1003089.
    NT2RM4000086, NT2RM4000265, NT2RM4000414, NT2RM4000779, NT2RM4000855, NT2RM4001160,
    NT2RM4001313, NT2RM4001437, NT2RM4001754, NT2RM4001953, NT2RM4001984, NT2RP200007 7,
    NT2RP2000183, NT2RP2000420, NT2RP2000678, NT2RP2000715, NT2RP2000842, NT2RP20009 70,
    NT2RP2001149,
                  NT2RP2001226.
                                NT2RP2001295, NT2RP2001347, NT2RP2001569,
                                                                           NT2RP2001
    NT2RP2001936, NT2RP2002041,
                                NT2RP2002172, NT2RP2002219, NT2RP2002316,
                                                                           NT2RP200 2546,
    NT2RP2002591, NT2RP2002643,
                                NT2RP2002741, NT2RP2002750, NT2RP2002778,
                                                                           NT2RP20 02857.
    NT2RP2003000, NT2RP2003073,
                                NT2RP2003237, NT2RP2003394, NT2RP2003517,
                                                                           NT2RP2 003668.
    NT2RP2003988, NT2RP2004232,
                                NT2RP2004523, NT2RP2004736, NT2RP2004767.
                                                                           NT2RP 2004775.
    NT2RP2004961, NT2RP2004962,
                                NT2RP2004982, NT2RP2005407, NT2RP2005726,
                                                                           NT2R P2006258.
    NT2RP2006261, NT2RP2006454,
                                NT2RP3000055, NT2RP3000233, NT2RP3000341,
                                                                           NT2 RP3000418.
    NT2RP3000451, NT2RP3000561,
                                NT2RP3000582, NT2RP3001281, NT2RP3001339, NT 2RP3001340,
                                NT2RP3001580, NT2RP3001589, NT2RP3002004, N T2RP3002173,
    NT2RP3001383, NT2RP3001432,
    NT2RP3003133, NT2RP3003346,
                                NT2RP3003403, NT2RP3003576, NT2RP3003625, NT2RP3003665,
    NT2RP3003800, NT2RP3003828,
                                 NT2RP3004070, NT2RP3004470, NT2RP4000023,
                                                                             NT2RP4000035,
    NT2RP4000102, NT2RP4000167, NT2RP4000214, NT2RP4000218, NT2RP400042 4, NT2RP4000915,
    NT2RP4002075.
    OVARC1000085, OVARC1000092, OVARC1000145, OVARC1000414, OVARC1000496, OVARC1000526,
    OVARC1000948, OVARC1001011, OVARC1001600, OVARC1001805, OVARC1001813, OVARC100184 6,
25
    PLACE1000540, PLACE1000599, PLACE1001088, PLACE1001377, PLACE1001440,
                                                                             PLACE1001517.
    PLACE1001672, PLACE1001756,
                                PLACE1002157, PLACE1002205, PLACE1002259, PLACE100239 9.
    PLACE1002477, PLACE1002583.
                                PLACE1002968, PLACE1003238, PLACE1003566,
                                                                           PLACE10035 93.
    PLACE1003618, PLACE1004274,
                                PLACE1004716, PLACE1004773, PLACE1004815.
                                                                           PLACE1004 979.
    PLACE1005052, PLACE1005086,
                                PLACE1005128, PLACE1005176, PLACE1005467.
                                                                           PLACE100 5639.
    PLACE1005850, PLACE1006003,
                                                                           PLACE10 06629,
                                PLACE1006017, PLACE1006288, PLACE1006371,
                                PLACE1008584, PLACE1008851, PLACE1008941,
    PLACE1007478, PLACE1008330.
                                                                           PLACE1 009039,
    PLACE1009493, PLACE1009539.
                                PLACE1009637, PLACE1009947, PLACE1010231,
                                                                           PLACE 1010562,
    PLACE1010579. PLACE1010739.
                                PLACE1010802, PLACE1010896, PLACE1011032,
                                                                           PLAC E1011185,
                                PLACE1011520, PLACE1011567, PLACE1011719,
    PLACE1011452, PLACE1011465.
                                                                           PLA CE2000011,
    PLACE2000017, PLACE2000061,
                                PLACE2000187, PLACE2000216, PLACE2000335,
                                                                           PL ACE2000347,
    PLACE2000366, PLACE2000394,
                                PLACE2000398, PLACE2000425, PLACE2000450, P LACE2000477,
    PLACE3000119,
                  PLACE3000207, PLACE3000230, PLACE3000271,
                                                              PLACE3000373.
                                                                             PLACE3000399.
    PLACE3000401, PLACE3000406, PLACE4000247, PLACE4000320, PLACE4000367, PLACE4000401,
    THYRO1000111, THYRO1000187, THYRO1000484, THYRO1000596, THYRO1000625, THYRO1000815,
    THYRO1000865, THYRO1001003, THYRO1001031, THYRO1001133, THYRO1001401, THYRO100142 6,
    THYRO1001434, THYRO1001559, THYRO1001570, THYRO1001706, THYRO1001746, THYRO10017 72,
    THYRO1001907, Y79AA1000033, Y79AA1000346, Y79AA1000805, Y79AA1001692, Y79AA1002 220,
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Group (5): 24 clones

45

The following 24 clones having low score in the ATGpr1 were judged to be full-length since their ATGpr2 score was 0.3 or higher (Table 11). The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA1000622, HEMBA1000749, HEMBA1000876, HEMBA1001226, HEMBA1001391, HEMBA1002742, HEMBA1003908, HEMBA1004000, HEMBB1000336, MAMMA1000356, MAMMA1000883, MAMMA100159 0, NT2RP2000289, NT2RP2001467, PLACE1002853, PLACE1003420, PLACE1004836, PLACE10049 13, PLACE1006795, PLACE1008181, PLACE1008715, PLACE4000344, THYRO1000129, THYRO1001 321,

Group (6): 65 clones

The following 65 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which at least either of the 5'-end or 3'-end sequence is not identical to any of human ESTs.

HEMBA1000604, HEMBA1000673, HEMBA1001024, HEMBA1001026, HEMBA1001734, HEMBA1001784,

HEMBA1001808, HEMBA1003902, HEMBA1004164, HEMBA1004909, HEMBA1005232, HEMBA100557 7, HEMBA1006461, HEMBA1006695, HEMBB1001119, HEMBB1001337, HEMBB1001536, HEMBB10018 68, HEMBB1002045, HEMBB1002579, MAMMA1000444, MAMMA1000761, MAMMA1000943, MAMMA1001 820, MAMMA1002360,

NT2RM4000366, NT2RM4001856, NT2RM4002390, NT2RP2000108, NT2RP2000257, NT2RP2001506, NT2RP2002047, NT2RP2002066, NT2RP2002475, NT2RP2004400, NT2RP2004587, NT2RP200528 9, NT2RP2005694, NT2RP3001898, NT2RP3003264, NT2RP3003433, NT2RP3003842, OVARC10012 40, PLACE1001323, PLACE1002227, PLACE1002500, PLACE1002604, PLACE1002772, PLACE1003 478, PLACE1004681, PLACE1005108, PLACE1005932, PLACE1006318, PLACE1006368, PLACE100 6506, PLACE1006904, PLACE1007557, PLACE1007877, PLACE1009048, PLACE1011109, PLACE10 11643, PLACE4000548, THYRO1000279, Y79AA1000410, Y79AA1002103,

Group (7): 32 clones

The following 32 clones having low scores in both the ATGpr1 and ATGpr2 (Table 11) were judged to be full-length, since both scores were still the maximum in a cluster compared with those of the other clones(at least 2 clones or more) in the same cluster. The clones were novel clones, in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5.

HEMBA100251, HEMBA1001803, HEMBA1001918, HEMBA1002257, HEMBA1003064, HEMBA1003714, HEMBA1004405, HEMBA1005508, HEMBB1000054, HEMBB1001142, MAMMA1000175, MAMMA100116 2, MAMMA1002972, NT2RM4000425, NT2RP2004512, NT2RP2005531, NT2RP2005942, NT2RP20065 54, NT2RP3001007, NT2RP3001318, OVARC1000017, OVARC1000068, OVARC1000486, PLACE1001 705, PLACE1002319, PLACE1007743, PLACE1007829, PLACE1008630, PLACE1009925, PLACE101 1492, PLACE1011749, THYRO1000793,

The following 117 clones, selected by assembling the sequence of the other clones in the same cluster and human ESTs, have high fullness ratio. The clones were classified into the following group (8) and (9). HEMBA1001323, HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1002538, HEMBA1003309, HEMBA1003939, HEMBA1004015, HEMBA1004295, HEMBA100467 2, HEMBA1004865, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1006779, HEMBA10072 88, HEMBB1000218, HEMBB1000272, HEMBB1000399, HEMBB1000491, HEMBB1000996, HEMBB1001 114. HEMBB1001850, HEMBB1002015, MAMMA1000287, MAMMA1001683, MAMMA1001686, MAMMA100 2612, NT2RM2000609, NT2RM4002438, NT2RM4002567, NT2RP2000270, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002736, NT2RP2002753, NT2RP2003456, NT2RP200372 7, NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2004580, NT2RP20052 93, NT2RP2005476, NT2RP2005753, NT2RP2005815, NT2RP2005841, NT2RP2005857, NT2RP2006 393, NT2RP2006467, NT2RP3000109, NT2RP3000449, NT2RP3001245, NT2RP3001634, NT2RP300 2056, NT2RP3002810, NT2RP3002955, NT2RP3003032, NT2RP3003138, NT2RP3003500, NT2RP30 03819, NT2RP400078, NT2RP4000515, NT2RP4000517, NT2RP4001407, NT2RP4001889, NT2RP4 002905, OVARC1000071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691, PLACE1001746, PLACE1001748, PLACE1001845, PLACE1002066, PLACE1003373, PLACE100390 0, PLACE1004118, PLACE1004256, PLACE1004284, PLACE1004336, PLACE1004506, PLACE10049 34, PLACE1005077, PLACE1005111, PLACE1005409, PLACE1005730, PLACE1006076, PLACE1006 360, PLACE1006470, PLACE1006760, PLACE1006867, PLACE1007045, PLACE1007111, PLACE100 7807,

Group (8): 36 clones

PLACE1008080, PLACE1008244, PLACE1008369,

THYRO1001365, THYRO1001673, Y79AA1001848,

Among the 117 clones described above, the following 36 clones were novel clones, in which at least either of the 5'-end or 3'-end sequence, or both of them were not identical to any of human ESTs.

PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, PLACE3000276, THYRO1 000805,

PLACE1008405, PLACE1008426,

PLACE10 08621,

HEMBA1001330, HEMBA1001712, HEMBA1001820, HEMBA1002204, HEMBA1002349, HEMBA1003939, HEMBA1004295, HEMBA1004865, HEMBA1006779, HEMBB1000218, HEMBB1000491, HEMBB100099 6, HEMBB1001114, MAMMA1000287, MAMMA1001686, MAMMA1002612, NT2RP2002736, NT2RP20037 27, NT2RP2004580, NT2RP2005841, NT2RP2006393, PLACE1002066, PLACE1003373, PLACE1003 900, PLACE1004118, PLACE1004336, PLACE1004506, PLACE1004934, PLACE1005409, PLACE100 5730, PLACE1006470, PLACE1008080, PLACE3000276, THYRO1000805, THYRO1001365, THYRO10 01673,

Group (9): 81 clones

Among the 117 clones described above, the following 81 clones were the clones in which the number of the identical

human ESTs for the 5'-end sequence is not more than 20 (except clones in which at least either of the 5'-end or 3'-end sequence, or both of them are not identical to any of human ESTs, and clones in which the number of the identical human ESTs for both the 5'-end and 3'-end sequences is 1 or higher and not more than 5).

HEMBA1001323, HEMBA1002538, HEMBA1003309, HEMBA1004015, HEMBA1004672, HEMBA1005251, HEMBA1006158, HEMBA1006676, HEMBA1007288, HEMBB1000272, HEMBB1000399, HEMBB100185 0, HEMBB1002015, MAMMA1001683, NT2RM2000609, NT2RM4002438, NT2RM4002567, NT2RP20002 70, NT2RP2000758, NT2RP2001290, NT2RP2001526, NT2RP2002124, NT2RP2002753, NT2RP2003 456, NT2RP2003871, NT2RP2003968, NT2RP2004321, NT2RP2004412, NT2RP2005293, NT2RP200 5476, NT2RP2005753, NT2RP2005815, NT2RP2005857, NT2RP2006467, NT2RP3000109, NT2RP30 00449, NT2RP3001245, NT2RP3001634, NT2RP3002056, NT2RP3002810, NT2RP3002955, NT2RP3 003032, NT2RP3003138, NT2RP3003500, NT2RP3003819, NT2RP4000078, NT2RP4000515, NT2RP 4000517,

NT2RP4001407, NT2RP4001889, NT2RP4002905,
OVARC1000071, OVARC1001883, PLACE1000292, PLACE1001007, PLACE1001395, PLACE1001691,
PLACE1001746, PLACE1001748, PLACE1001845, PLACE1004256, PLACE1004284, PLACE100507 7,
PLACE1005111, PLACE1006076, PLACE1006360, PLACE1006760, PLACE1006867, PLACE10070 45,
PLACE1007111, PLACE1007807, PLACE1008244, PLACE1008369, PLACE1008405, PLACE1008 426,
PLACE1008621, PLACE1009020, PLACE1009621, PLACE1010089, PLACE1010270, Y79AA100 1848,

Group (10): 938 clones

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The following 938 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 4. The clones were novel clones, in which at least the 5'-end sequence was not identical to any of human ESTs.

HEMBA100042, HEMBA1000213, HEMBA1000264, HEMBA1000355, HEMBA1000366, HEMBA1000411, HEMBA1000422, HEMBA1000428, HEMBA1000434, HEMBA1000442, HEMBA1000442, HEMBA1000442, HEMBA1000442, HEMBA1000702, HEMBA1000747, HEMBA1000769, HEMBA1000773, HEMBA1000744, HEMBA1000774, HEMBA1000852, HEMBA1000854, HEMBA1000908, HEMBA1000985, HEMBA1000985, HEMBA1001017, HEMBA1001019, HEMBA1001094, HEMBA1001121, HEMBA1001172, HEMBA1001172, HEMBA1001327, HEMBA1001557, HEMBA1001566, HEMBA1001585, HEMBA1001585, HEMBA1001585, HEMBA1001678, HEMBA1001678, HEMBA1001681, HEMBA1001718, HEMBA1001761, HEMBA1001945, HEMBA1001960, HEMBA1001964, HEMBA1001991.

HEMBA1002003, HEMBA1002008, HEMBA1002039, HEMBA1002100, HEMBA1002113, HEMBA1002139, HEMBA1002144, HEMBA1002153, HEMBA1002226, HEMBA1002253, HEMBA1002270, HEMBA100232 1, HEMBA1002558, HEMBA1002551, HEMBA1002551, HEMBA1002552, HEMBA1002552, HEMBA1002552, HEMBA1002553, HEMBA1002666, HEMBA1002666, HEMBA1002678, HEMBA1002678, HEMBA1002712, HEMBA1002712, HEMBA1002714, HEMBA1002748, HEMBA1002780, HEMBA1002801, HEMBA1002826, HEMBA1002833, HEMBA1002896, HEMBA1003033, HEMBA1003033, HEMBA10030334, HEMBA1003034, HEMBA1003334, HEMBA1003334, HEMBA1003336, HEMBA1003336, HEMBA1003376, HEMBA1003376, HEMBA1003758, HEMBA1003773, HEMBA1003784, HEMBA1003885, HEMBA1003937, HEMBA1003950,

HEMBA1003976,
HEMBA1004012, HEMBA1004045, HEMBA1004138, HEMBA1004241, HEMBA1004272, HEMBA1004306, HEMBA1004312, HEMBA1004334, HEMBA1004354, HEMBA1004356, HEMBA1004366, HEMBA100439 4, HEMBA1004366, HEMBA1004429, HEMBA1004460, HEMBA1004617, HEMBA1004670, HEMBA1004733, HEMBA1004816, HEMBA1004820, HEMBA1004918, HEMBA1004960, HEMBA1004978, HEMBA1004995, HEMBA1005008, HEMBA1005062, HEMBA1005293, HEMBA10050304, HEMBA1005315, HEMBA1005372, HEMBA1005370, HEMBA1005616, HEMBA1005627, HEMBA1005631, HEMBA1005632, HEMBA1005631, HEMBA1005632, HEMBA1005630, HEMBA1005631, HEMBA1005632, HEMBA1005699, HEMBA1005705, HEMBA1005765, HEMBA1005780, HEMBA1005853, HEMBA1005909,

HEMBA1005931, HEMBA1005934, HEMBA1005962, HEMBA1005999,
HEMBA1006002, HEMBA1006005, HEMBA1006042, HEMBA1006067, HEMBA1006090, HEMBA1006142,
HEMBA1006268, HEMBA1006334, HEMBA1006359, HEMBA1006416, HEMBA1006419, HEMBA100642 1,
HEMBA1006446, HEMBA1006471, HEMBA1006486, HEMBA1006489, HEMBA1006546, HEMBA10065 69,

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HEMBA1006595, HEMBA1006617, HEMBA1006635, HEMBA1006648, HEMBA1006665, HEMBA1006 821,
HEMBA1006849, HEMBA1006865, HEMBA1006929, HEMBA1007045, HEMBA1007051, HEMBA100 7073.
HEMBA1007113, HEMBA1007256, HEMBA1007273, HEMBA1007322, HEMBA1007347, HEMBB1000039.
HEMBB1000044, HEMBB1000059, HEMBB1000089, HEMBB1000099, HEMBB1000113, HEMBB1000141,
HEMBB1000250, HEMBB1000264, HEMBB1000274, HEMBB1000284, HEMBB100030 7, HEMBB1000312.
HEMBB1000317, HEMBB1000337, HEMBB1000341, HEMBB1000343, HEMBB10003 69, HEMBB1000376,
HEMBB1000434. HEMBB1000441, HEMBB1000455, HEMBB1000472, HEMBB1000 518, HEMBB1000530,
HEMBB1000564, HEMBB1000575, HEMBB1000586, HEMBB1000598, HEMBB100 0637, HEMBB1000638,
HEMBB1000643, HEMBB1000665, HEMBB1000726, HEMBB1000738, HEMBB10 00770, HEMBB1000794,
HEMBB1000821, HEMBB1000822, HEMBB1000883, HEMBB1000888, HEMBB1 000890, HEMBB1000910,
HEMBB1000959, HEMBB1000981, HEMBB1001037, HEMBB1001051, HEMBB 1001060, HEMBB1001133,
HEMBB1001208, HEMBB1001209, HEMBB1001218, HEMBB1001221, HEMB B1001249, HEMBB1001254.
HEMBB1001267, HEMBB1001282, HEMBB1001302, HEMBB1001304, HEM BB1001335, HEMBB1001348,
HEMBB1001356, HEMBB1001366, HEMBB1001380, HEMBB1001443, HE MBB1001463, HEMBB1001464.
HEMBB1001521, HEMBB1001527, HEMBB1001537, HEMBB1001555, HEMBB1001586, HEMBB1001619,
HEMBB1001630, HEMBB1001637, HEMBB1001641, HEMBB1001706, HEMBB1001735, HEMBB100175 3,
HEMBB1001762, HEMBB1001802, HEMBB1001880, HEMBB1001899, HEMBB1001921, HEMBB10019 30,
HEMBB1001952, HEMBB1001953, HEMBB1001973, HEMBB1001988, HEMBB1002002, HEMBB1002 005,
HEMBB1002009, HEMBB1002042, HEMBB1002049, HEMBB1002069, HEMBB1002115, HEMBB100 2139,
HEMBB1002189, HEMBB1002232, HEMBB1002254, HEMBB1002255, HEMBB1002280, HEMBB10 02306,
HEMBB1002364, HEMBB1002371, HEMBB1002381, HEMBB1002453, HEMBB1002477, HEMBB1 002509,
HEMBB1002531, HEMBB1002601, HEMBB1002603, HEMBB1002610, HEMBB1002613, HEMBB 1002623,
HEMBB1002635, HEMBB1002664, HEMBB1002677, HEMBB1002683, HEMBB1002686, HEMB B1002699.
HEMBB1002712.
MAMMA1000009, MAMMA1000045, MAMMA1000103, MAMMA1000133, MAMMA1000134, MAMMA1000155,
MAMMA1000241, MAMMA1000254, MAMMA1000264, MAMMA1000266, MAMMA1000340, MAMMA100038 5,
MAMMA1000423, MAMMA1000424, MAMMA1000446, MAMMA1000478, MAMMA1000501, MAMMA10005 16,
MAMMA1000565, MAMMA1000585, MAMMA1000605, MAMMA1000616, MAMMA1000621, MAMMA1000 643.
MAMMA1000669, MAMMA1000696, MAMMA1000707, MAMMA1000714, MAMMA1000718, MAMMA100 0732.
MAMMA1000733, MAMMA1000839, MAMMA1000845, MAMMA1000855, MAMMA1000863, MAMMA10 00867.
MAMMA1000880, MAMMA1000905, MAMMA1000942, MAMMA1000957, MAMMA1000987, MAMMA1 000998.
MAMMA1001003, MAMMA1001024, MAMMA1001030, MAMMA1001035, MAMMA1001050, MAMMA 1001067.
MAMMA1001082, MAMMA1001092, MAMMA1001145, MAMMA1001161, MAMMA1001203, MAMM A1001243,
MAMMA1001268, MAMMA1001280, MAMMA1001298, MAMMA1001322, MAMMA1001324, MAM MA1001383,
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Group (11): 228 clones

MAMMA1003140.

The following 228 clones having low scores in both the ATGpr1 and ATGpr2 were judged to be full-length according to the fullness ratio shown in Table 7. The clones were a novel clone, in which at least the 3'-end sequence was not identical to any of human ESTs.

HEMBA1000111, HEMBA1000243, HEMBA1000338, HEMBA1000390, HEMBA1000418, HEMBA1000459, HEMBA1000518, HEMBA1000791, HEMBA1000974, HEMBA1001007, HEMBA1001077, HEMBA100109 9, HEMBA1001123, HEMBA1001294, HEMBA1001299, HEMBA1001442, HEMBA1001861, HEMBA10019 42, HEMBA1002592, HEMBA1002679, HEMBA1002944, HEMBA1003034, HEMBA1003142, HEMBA1003571, HEMBA1003579, HEMBA1003598, HEMBA1003034, HEMBA1003958, HEMBA1003034, HEMBA1003034, HEMBA1003035, HEMBA1003034, HEMBA1003035, HEMBA1003038, HEMBA1003039, HEMBA1005497, HEMBA1005552, HEMBA1005359, HEMBA1005717, HEMBA1005829, HEMBA1006938, HEMBA1006328, HEMBA1006334, HEMBA1006540, HEMBA1006780, HEMBA1006921, HEMBA1006938, HEMBA1007017, HEMBA1007341,

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EXAMPLE 12

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Homology search using the 5'-end and 3'-end sequences of the selected clones.

[0172] The 5' -end sequences of the selected 5547 clones were used for the homology search of the SwissProt, and both the 5' -end 3' -end sequences were used for the search of the GenBank and UniGene (ref. the result of the search of the SwissProt, GenBank (except ESTs and STSs), and UniGene (Human) was attached).

[0173] Each search result is shown in the last part of this SPECIFICATION by arranging each item in the following format.

		5'-end sequence	3'-end sequence
	Swiss-Prot	Homology search result 1	
	GenBank	Homology search result 2	Homology search result 3
15	UniGene	Homology search result 4	Homology search result 5

[0174] According to the top hit data, at least 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth. Among the clones predicted belonging to any of the categories, 1001 clones were estimated to have a relatively high homology with the known proteins or genes in the same category. In addition, 429 clones were estimated to have a relatively low homology with the known proteins in the same category. [0175] Herein, the term "relatively high homology" is defined as having 60% or more identity and the P-value 10-10 or less in comparison with known sequences in the SwissProt database, or 64% or more identity and the P-value 10⁻¹⁵ or less in comparison with those in the GenBank and UniGene databases (see the attached list). Also, the term "relatively low homology" is defined as not fulfilling the requirements to be "relatively high homology", but still having the scores, 25% or more identity and the P-value 10-6 or less, using the sequence having 55 nucleotides or more, in comparison with known sequences in the SwissProt database (see the attached list). The P-value is a score obtained statistically by taking into account the probability of occurrence of the similarity between two sequences. In general, the smaller P-value reflects the higher similarity (Altschul S.F., Gish W., Miller W., Myers E.W., and Lipman D.J. (1990) "Basic local alignment search tool" J.Mol. Biol., 215: 403-410; Gish W., and States D.J. (1993) "Identification of protein coding regions by database similarity search" Nature Genet. 3: 266-272).

[0176] The clones predicted to encode a protein in the category of secretory protein or membrane protein have the keywords, "signal", "transmembrane", "membrane", "extracellular matrix", "receptor", "G-protein coupled receptor", "ionic channel", "voltage-gated channel", "calcium channel", "cell adhesion", "collagen", or "connective tissue", or descriptions from which the clone can be predicted to be a secretory or membrane protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0177] The clones predicted to encode a protein in the category of glycoprotein have the keywords, "glycoprotein", or descriptions from which the clone can be predicted to be a glycoprotein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence. [0178] The clones predicted to encode a protein in the category of proteins associated with signal transduction have the keywords, "serine/threonine-protein kinase", "tyrosine-protein kinase", "SH3 domain", or "WD repeat", or descriptions from which the clone can be predicted to be a protein associated with signal transduction (such as "ADP-ribosylation factor"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0179] The clones predicted to encode a protein in the category of proteins associated with transcription have the keywords, "transcription regulation", "zinc finger", or "homeobox", or descriptions from which the clone can be predicted to be a protein associated with transcription, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0180] The clones predicted to encode a protein in the category of proteins associated with diseases are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein that is deposited in the Online Mendelian Inheritance in Man (OMIM) database, which is a database of human genes and diseases, or the top hit data has descriptions from which the clone can be predicted to be a protein associated with diseases.

[0181] The clones predicted to encode a protein in the category of enzyme or proteins associated with metabolism are the clones in which the top hit data of the SwissProt using the 5'-end sequence, or the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence is a gene or protein with E.C.No. (Enzyme commission number), or the top hit data has descriptions from which the clone can be predicted to be an enzyme or protein associated with metabolism (such as "metabolism", "oxidoreductase", or "lipid").

[0182] The clones predicted to encode a protein in the category of proteins associated with cell division or cell proliferation have the keywords, "cell division", "cell cycle", "mitosis", or "chromosomal protein", or descriptions from which the clone can be predicted to be a protein associated with cell division or cell proliferation (such as "histone", "cell growth", or "apoptosis"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0183] The clones predicted to encode a protein in the category of proteins associated with cytoskeleton have the keywords, "structural protein", "cytoskeleton", "actin-binding", or "microtubules", or descriptions from which the clone can be predicted to be a protein associated with cytoskeleton, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0184] The clones predicted to encode a protein in the category of proteins associated with RNA synthesis include the above clones predicted to be a protein associated with transcription, and also the clones which have the keywords, "RNA splicing", or "RNA processing", or descriptions from which the clone can be predicted to be a protein associated with RNA synthesis (such as "RNA helicase", "polyadenylation", or "RNA transport"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

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[0185] The clones predicted to encode a protein in the category of nuclear protein include the above clones predicted to be a protein associated with transcription, and also the clones which have the keyword, "nuclear protein", or descriptions from which the clone can be predicted to be a nuclear protein, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0186] The clones predicted to encode a protein in the category of proteins associated with protein synthesis or transport have the keywords, "translation regulation", "protein biosynthesis", "amino-acid biosynthesis", "ribosomal protein", or "protein transport", or descriptions from which the clone can be predicted to be a protein associated with protein synthesis or transport (such as "signal recognition particle", "ubiquitin", "proteosome", or "protease"), in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0187] The clones predicted to encode a protein in the category of proteins associated with cellular defense have the keywords, "heat shock", "chaperone", "DNA repair", or "DNA damage", or descriptions from which the clone can be predicted to be a protein associated with cellular defense, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0188] The clones predicted to encode a protein in the category of proteins associated with development or growth have the keyword, "developmental protein", or descriptions from which the clone can be predicted to be a protein associated with development or growth, in the top hit data of the SwissProt using the 5'-end sequence, or in the top hit data of the GenBank or UniGene using the 5'-end sequence and 3'-end sequence.

[0189] The following 1430 clones were predicted to encode a protein belonging to any of the categories, secretory or membrane protein, glycoprotein, protein associated with signal transduction, protein associated with transcription, protein associated with diseases, enzyme or protein associated with metabolism, protein associated with cell division or cell proliferation, protein associated with cytoskeleton, protein associated with RNA synthesis, nuclear protein, protein associated with protein synthesis or transport, protein associated with cellular defense, or protein associated with development or growth.

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[0190] Among the clones, 1001 clones were estimated to have a relatively high homology with the known proteins
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     [0191] The following 429 clones were estimated to have a relatively low homology with the known proteins in the
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55

same category.

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10		НЕМВА1002102, НЕМВА1002609, НЕ		, HEMBA1002212, EMBA1002939,	HEMBA1002495,
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	HEMBA1003071, HEMBA1003291,	,	5,
5	HEMBA1003568, HEMBA1003953,	HEMBA1003598, HEMBA1003680, HEMBA1003684, HEMBA1003830	5,
10	HEMBA1004276,	HEMBA1004408, HEMBA1004669, HEMBA1004734,	
10	HEMBA1005475,	HEMBA1005595, HEMBA1005737, HEMBA1005815, HEMBA1005931,	
15	HEMBA1006347, HEMBA1006737,	HEMBA1006474, HEMBA1006521, HEMBA1006562, HEMBA1006652 HEMBA1006914,	2,
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30	MAMMA1002143, MAMMA1002671,	MAMMA1002573, MAMMA1002619, MAMMA1002636, MAMMA1002650 MAMMA1002881, MAMMA1002937,),
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45	NT2RP3000994,		·	•	•
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	(1) Among the 1430 clones, the following 259 clones were predicted to encode a protein in the category of secretory protein or membrane protein (including the clones belonging to plural categories). HEMBA1000005, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001351, HEMBA1001407,					
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40		HEMBA1002688, HEMI HEMBA1003598, HEMI				
		HEMBA1005338, HEMI				
		HEMBA1006310, HEM				

HEMBA1007301, HEMBB1000037, HEMBB1000119, HEMBB1000317, HEMBB1000530, HEMBB1 000915, 45 HEMBB1001112, HEMBB1001119, HEMBB1001151, HEMBB1001282, HEMBB1001331, HEMBB 1002042, HEMBB1002092, HEMBB1002193, HEMBB1002600, MAMMA1000103, MAMMA1000424, MAMM A1000672, MAMMA1000734, MAMMA1000897, MAMMA1001008, MAMMA1001030, MAMMA1001041, MAM MA1001075, MAMMA1001080, MAMMA1001751, MAMMA1001754, MAMMA1002143, MAMMA1002219, MA MMA1002485, MAMMA1002636, MAMMA1002938, MAMMA1003104, NT2RM1000199, NT2RM1000355, N T2RM1000430, NT2RM1000648, NT2RM1000742, NT2RM1000770, NT2RM1000811, NT2RM1000833, NT2RM1000867, NT2RM1000882, NT2RM1000883, NT2RM1001139, NT2RM2000374, NT2RM2000402, NT2RM2000422, NT2RM2000566, NT2RM2000623, NT2RR2001324, NT2RM2001499, NT2RM200158 8, NT2RM2001592, NT2RM2001613, NT2RM2001635, NT2RM2001648, NT2RM2001760, NT2RM20021 09, NT2RM2002145, NT2RM4000156, NT2RM4000169, NT2RM4000386, NT2RM4000433, NT2RM4000795, NT2RM4001032, NT2RM4001054, NT2RM4001410, NT2RM4001605; NT2RM4001930, NT2RM4001987, NT2RM4002073, NT2RM4002145, NT2RM4002189, NT2RM4002558, NT2RM400256 5, NT2RM4002571, NT2RM4002594, NT2RP1000130, NT2RP1000326, NT2RP1000363, NT2RP10004 13, NT2RP1000547, NT2RP1000782, NT2RP1000856, NT2RP1001247, NT2RP1001313, NT2RP1001 546, NT2RP1001569, NT2RP2000056,

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NT2RP2000257, NT2RP2000842, NT2RP2001081, NT2RP200 1991, NT2RP2002025, NT2RP2002193,
       NT2RP2002256, NT2RP2002312, NT2RP2002479, NT2RP20 02504, NT2RP2003433, NT2RP2003706,
       NT2RP2003760, NT2RP2004194, NT2RP2004655, NT2RP2 005020, NT2RP2005344, NT2RP2005465,
       NT2RP2005752, NT2RP2005753, NT2RP2005835, NT2RP 2005933, NT2RP2006275, NT2RP2006565,
       NT2RP2006571, NT2RP3000324, NT2RP3000739, NT2R P3000919, NT2RP3001140, NT2RP3001355,
       NT2RP3001676, NT2RP3001708, NT2RP3001727, NT2 RP3002659, NT2RP3003059, NT2RP3003101,
       NT2RP3003203, NT2RP3003212, NT2RP3003242, NT 2RP3003672, NT2RP3003701, NT2RP3003716,
       NT2RP3003918, NT2RP3004207, NT2RP3004253, N T2RP4000008, NT2RP4000212, NT2RP4000243,
       NT2RP4000417, NT2RP4000878, NT2RP4000907, NT2RP4000925, NT2RP4000928, NT2RP4001079,
       NT2RP4001117, NT2RP4001150, NT2RP4001174, NT2RP4001313, NT2RP4001315, NT2RP4001345,
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       NT2RP4001372, NT2RP4001574, NT2RP400182 3, NT2RP4001966, NT2RP5003524, OVARC1000060,
       OVARC1000326, OVARC1000543, OVARC10006 82, OVARC1000722, OVARC1000912, OVARC1001055,
       OVARC1001154, OVARC1001261,
       OVARC1001329, OVARC1001419, OVARC1001506, OVARC1001542, OVARC1001610, OVARC1001713,
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       OVARC1001802, OVARC1001943, PLACE1000611, PLACE1001092, PLACE1001136, PLACE100129 4,
       PLACE1002437, PLACE1002500, PLACE1002722, PLACE1002782, PLACE1002816, PLACE10034 20,
       PLACE1003493, PLACE1003596, PLACE1004104, PLACE1004197, PLACE1004203, PLACE1004 277,
       PLACE1004629, PLACE1004751, PLACE1005494, PLACE1005804, PLACE1005890, PLACE100 6157,
       PLACE1007409, PLACE1007852, PLACE1008044, PLACE1008273, PLACE1008309, PLACE10 08603,
20
       PLACE1008643, PLACE1009659, PLACE1009670, PLACE1009845, PLACE1009861, PLACE1 011371,
       PLACE1011492, PLACE1011896, PLACE2000034, PLACE2000341, PLACE2000399, PLACE 2000458,
       PLACE3000020, PLACE3000322, PLACE3000353, PLACE4000052, PLACE4000269, PLAC E4000522,
       PLACE4000581, THYRO1000783, Y79AA1000258, Y79AA1000346, Y79AA1001874,
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25 (2) The following 111 clones were predicted to encode a protein in the category of glycoprotein (including the clones belonging to plural categories). HEMBA1000752, HEMBA1001017, HEMBA1001059, HEMBA1001071, HEMBA1001302, HEMBA1002150, HEMBA1002547, HEMBA1002555, HEMBA1003077, HEMBA1003538, HEMBA1003598, HEMBA100386 6, HEMBA1004356, HEMBA1005338, HEMBA1005576, HEMBA1005699, HEMBA1006562, HEMBA10069 76, 30 HEMBA1007301, HEMBB1000317, HEMBB1000530, HEMBB1001119, HEMBB1002092, HEMBB1002 193, MAMMA1000672, MAMMA1000897, MAMMA1001030, MAMMA1001388, MAMMA1001771, MAMMA100 2485, MAMMA1002636, MAMMA1002938, NT2RM1000648, NT2RM2000374, NT2RM2000407, NT2RM20 00422, NT2RM2000623, NT2RM2001499, NT2RM2001588, NT2RM2001839, NT2RM2001930, NT2RM4 000156, NT2RM4000233, NT2RM4001410, NT2RM4001987, NT2RM4002145, NT2RM4002189, NT2RM 4002194, 35 NT2RM4002558, NT2RM4002571, NT2RP1000856, NT2RP1001247, NT2RP2000056, NT2R P2001991, NT2RP2002025, NT2RP2002325, NT2RP2002385, NT2RP2003706, NT2RP2005933, NT2 RP3001140, NT2RP3003242, NT2RP3003672, NT2RP3003701, NT2RP3003716, NT2RP3003914, NT 2RP3004253, NT2RP4000212, NT2RP4000417, NT2RP4000925, NT2RP4001149, NT2RP4001150, N T2RP4001219, NT2RP4001345, NT2RP4001372, NT2RP5003522, NT2RP5003524, OVARC1000543, OVARC1000682, OVARC1000722, OVARC1001055, OVARC1001154, OVARC1001506, OVARC1001713 PLACE1002437, 40 PLACE1002722, PLACE1003258, PLACE1003493, PLACE1004197, PLACE100475 1, PLACE1005698,

(3) The following 147 clones were predicted to encode a protein in the category of proteins associated with signal transduction (including the clones belonging to plural categories).

PLACE4000581, THYRO1000852, Y79AA1001874,

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PLACE1005804, PLACE1006157, PLACE1006754, PLACE1008643, PLACE10090 94, PLACE1009861, PLACE1011371, PLACE2000034, PLACE2000062, PLACE2000399, PLACE2000458, PLACE3000099, PLACE3000020, PLACE3000353, PLACE3000373, PLACE4000522, PLACE4000520, PLACE4000522,

HEMBA100030, HEMBA1000185, HEMBA1000303, HEMBA1000459, HEMBA1000491, HEMBA1000657, HEMBA1000991, HEMBA1001174, HEMBA1001387, HEMBA1001822, HEMBA1001921, HEMBA100221 2, HEMBA1002341, HEMBA1002417, HEMBA1002810, HEMBA1002896, HEMBA1003291, HEMBA10033 14, HEMBA1003560, HEMBA1003615, HEMBA1003621, HEMBA1003805, HEMBA1004286, HEMBA1004 604, HEMBA1006173, HEMBA1006540, HEMBA1006973, HEMBA1007174, HEMBB1000632, HEMBB100 0781, HEMBB1001051, HEMBB1001177, HEMBB1001234, HEMBB1001294, HEMBB1001384, HEMBB10 02193, HEMBB1002477, HEMBB1002635, MAMMA1001305, MAMMA1002699, MAMMA1002842, MAMMA1 003057, MAMMA1003113, NT2RM1000118, NT2RM1000186, NT2RM1000772, NT2RM1000850, NT2RM 1001072, NT2RM2000124, NT2RM2000612, NT2RM2001065, NT2RM2001221, NT2RM2001345, NT2R M2001652, NT2RM2001983, NT2RM2002109, NT2RM2002128, NT2RM4000327, NT2RM4000354, NT2 RM4000798,

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NT2RM4001203, NT2RM4001411, NT2RM4001412, NT2RM4001582, NT2RM4001629, NT 2RM4001758, NT2RM4002226, NT2RP100018, NT2RP1000112, NT2RP1000701, NT2RP1000825, N T2RP1001395, NT2RP1001410, NT2RP1001457, NT2RP1001482, NT2RP2002058, NT2RP2002710, NT2RP2003702, NT2RP2003912, NT2RP2004232, NT2RP2004768, NT2RP2005022, NT2RP2005620, NT2RP2005890, NT2RP3000299, NT2RP3000403, NT2RP3000742, NT2RP3000845, NT2RP300176 4, NT2RP3002004, NT2RP3002785, NT2RP3002909, NT2RP3003230, NT2RP3003411, NT2RP30034 64, NT2RP3003800, NT2RP3004475, NT2RP3004534, NT2RP400147, NT2RP4000376, NT2RP4001122, NT2RP4001375, NT2RP4001644, NT2RP4001725, NT2RP4001760, NT2RP4001849, NT2RP4001927, NT2RP4002408, NT2RP5003477, OVARC1000437, OVARC1000465, OVARC100064 9, OVARC1000945, OVARC1001476, OVARC1001861, PLACE1001377, PLACE1001384, PLACE10013 87, PLACE1002493, PLACE1002591, PLACE1002685, PLACE1003353, PLACE1004128, PLACE1004 918, PLACE1005502, PLACE1005739, PLACE1005968, PLACE1006385, PLACE1007238, PLACE100 7375, PLACE1007488, PLACE1008244, PLACE1008650, PLACE1009319, PLACE1009468, PLACE10 10083, PLACE1010942, PLACE1011041, PLACE1011046, PLACE1011923, PLACE2000164, PLACE3 000145, PLACE3000350, PLACE3000475, THYRO1001595, Y79AA1000059, Y79AA1000966,
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(4) The following 230 clones were predicted to encode a protein in the category of proteins associated with transcription (including the clones belonging to plural categories). HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000851, HEMBA1001137,

HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA100184 7, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA1003545, HEMBA10035 68, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004097, HEMBA1004321, HEMBA1004 353, HEMBA1004479, HEMBA1004758, HEMBA1005359, HEMBA1005528, HEMBA1005548, HEMBA100 5931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006559, HEMBB1000250, HEMBB10 00789, HEMBB1001011, HEMBB1001314, HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1 002217. MAMMA1000155, MAMMA1000183, MAMMA1000388, MAMMA1001633, MAMMA1001820, MAMMA 1001837, MAMMA1002650, MAMMA1002937, NT2RM1000055, NT2RM1000666, NT2RM1000691, NT2R M1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2 RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2002580, NT2RM4000024, NT 2RM4000104. NT2RM4000191, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, N T2RM4000996. NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM4001746, NT2RM4001783. NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RP1000111, NT2RP1000333. NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP2000008, NT2RP200013 3, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2002252, NT2RP20025 03, NT2RP2002591, NT2RP2003125, NT2RP2003228, NT2RP2003293, NT2RP2003564,

NT2RP2003714, NT2RP2004013, NT2RP2004187, NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP2006238, NT2RP200646 4, NT2RP300050, NT2RP3000397, NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP30006 32, NT2RP3001057, NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001 399, NT2RP3001671, NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP300 1855, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP30 03251, NT2RP3003327, NT2RP3004258, NT2RP3004566, NT2RP3004572, NT2RP3004594, NT2RP3 004617, NT2RP4000078, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000787, NT2RP 4000865, NT2RP4000997, NT2RP4001029, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2R P4001551, NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP4002078, OVA RC1000479, OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577, OVARC1001949, PL ACE1000066. PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706, PLACE10011118, P LACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE1002450, PLACE1002532, PLACE1002834, PLACE1003302, PLACE1003605, PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004681, PLACE1006414, PLACE1006438, PLACE1006482, PLACE100723 9, PLACE1007346, PLACE1007544, PLACE1007598, PLACE1007955, PLACE1007969, PLACE10086 27, PLACE1008941, PLACE1009099, PLACE1009246, PLACE1009398, PLACE1010261,

PLACE1010491, PLACE1010702, PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072, PLACE2000359, PLACE4000128, PLACE4000156, PLACE4000259, PLACE4000431, THYRO100012 1, THYRO1000242, THYRO1000501, THYRO1000585, THYRO1001100, THYRO1001189, Y79AA10000 37, Y79AA1000627, Y79AA1000705, Y79AA1001105, Y79AA1001299, Y79AA1001391, Y79AA1001 533, Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1002093, Y79AA1002472, Y79AA100 2482,

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(5) The following 436 clones were predicted to encode a protein in the category of proteins associated with diseases
       (including the clones belonging to plural categories).
       HEMBA1000158, HEMBA1000216, HEMBA1000523, HEMBA1000561, HEMBA1000569, HEMBA1000673.
       HEMBA1000972, HEMBA1001017, HEMBA1001019, HEMBA1001059, HEMBA1001071, HEMBA100108 8,
       HEMBA1001123, HEMBA1001407, HEMBA1001569, HEMBA1001570, HEMBA1001672, HEMBA10020 35.
       HEMBA1002160, HEMBA1002162, HEMBA1002547, HEMBA1002939, HEMBA1003077, HEMBA1003 418,
       HEMBA1003433, HEMBA1003538, HEMBA1003555, HEMBA1003568, HEMBA1003569, HEMBA100 3581,
       HEMBA1003720, HEMBA1003760, HEMBA1004168, HEMBA1004248, HEMBA1004276, HEMBA10 04354,
       HEMBA1004356, HEMBA1004479, HEMBA1004534, HEMBA1004669, HEMBA1004752, HEMBA1 004753.
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       HEMBA1004807, HEMBA1005293, HEMBA1005331, HEMBA1005359, HEMBA1005367, HEMBA 1005423,
       HEMBA1005443, HEMBA1005475, HEMBA1005528, HEMBA1005576, HEMBA1005581, HEMB A1005679.
       HEMBA1005699, HEMBA1005732, HEMBA1005931, HEMBA1006173, HEMBA1006474, HEM BA1006562.
       HEMBA1006648, HEMBA1006652, HEMBA1006737, HEMBA1006807, HEMBA1006976, HE MBA1007078.
       HEMBA1007121, HEMBA1007243, HEMBA1007301, HEMBB1000264, HEMBB1000637, H EMBB1000684.
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       HEMBB1000693, HEMBB1000915, HEMBB1000927, HEMBB1001011, HEMBB1001051, HEMBB1001068.
       HEMBB1001102, HEMBB1001119, HEMBB1001133, HEMBB1001177, HEMBB1001802, HEMBB1001908,
       HEMBB1001922, HEMBB1001967, HEMBB1002050, HEMBB1002193, HEMBB100221 7, HEMBB1002358.
       HEMBB1002635, HEMBB1002683, MAMMA1000009, MAMMA1000092, MAMMA10001 03, MAMMA1000173,
       MAMMA1000731, MAMMA1000897, MAMMA1000956, MAMMA1001021,
       MAMMA1001041, MAMMA1001105, MAMMA1001198, MAMMA1001305, MAMMA1001576, MAMMA1001633,
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       MAMMA1002140, MAMMA1002170, MAMMA1002198, MAMMA1002268, MAMMA1002352, MAMMA100239 2,
       MAMMA1002485, MAMMA1002530, MAMMA1002573, MAMMA1002636, MAMMA1002858, MAMMA10028 69.
       MAMMA1002881, MAMMA1002937, MAMMA1003047, MAMMA1003099, MAMMA1003146, MAMMA1003 166,
       NT2RM1000035, NT2RM1000153, NT2RM1000257, NT2RM1000318, NT2RM1000377, NT2RM100 0394,
25
       NT2RM1000555, NT2RM1000691, NT2RM1000725, NT2RM1000770, NT2RM1000826, NT2RM10 00850,
       NT2RM1001059, NT2RM1001072, NT2RM1001082, NT2RM1001092, NT2RM2000363, NT2RM2 000374,
       NT2RM2000594, NT2RM2000599, NT2RM2000714, NT2RM2000740, NT2RM2000821, NT2RM 2001035,
       NT2RM2001499, NT2RM2001575, NT2RM2001605, NT2RM2001664, NT2RM2001670, NT2R M2001803,
       NT2RM2001823, NT2RM2001839, NT2RM4000085, NT2RM4000139, NT2RM4000155, NT2 RM4000290,
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       NT2RM4000689, NT2RM4000895, NT2RM4001320, NT2RM4001566, NT2RM4001629, NT 2RM4001741,
       NT2RM4001819, NT2RM4001823, NT2RM4001865, NT2RM4001940, NT2RM4002093, N T2RM4002146,
       NT2RM4002161, NT2RM4002189, NT2RM4002457, NT2RM4002558, NT2RM4002571, NT2RP1000086,
       NT2RP1000376, NT2RP1000574, NT2RP1000581, NT2RP1000629, NT2RP1000721, NT2RP1000733,
       NT2RP1000738, NT2RP1000825, NT2RP1000833, NT2RP1000860, NT2RP100096 6, NT2RP1001033,
       NT2RP1001247, NT2RP2000126, NT2RP2000147, NT2RP2000233, NT2RP20002 97, NT2RP2000310,
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       NT2RP2000414, NT2RP2000448, NT2RP2000459, NT2RP2000603,
       NT2RP2000812, NT2RP2001127, NT2RP2001327, NT2RP2001440, NT2RP2001520, NT2RP2001663,
       NT2RP2001748, NT2RP2001876, NT2RP2001898, NT2RP2002025, NT2RP2002066, NT2RP200225 9,
       NT2RP2002312, NT2RP2002503, NT2RP2002618, NT2RP2003228, NT2RP2003293, NT2RP20032 95,
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       NT2RP2003517, NT2RP2003564, NT2RP2003706, NT2RP2003737, NT2RP2003968, NT2RP2004 013,
       NT2RP2004098, NT2RP2004187, NT2RP2004316, NT2RP2004523, NT2RP2004538, NT2RP200 4933,
       NT2RP2005144, NT2RP2005288, NT2RP2005289, NT2RP2005325, NT2RP2005358, NT2RP20 05531,
       NT2RP2005773, NT2RP2006219, NT2RP3000050, NT2RP3000080, NT2RP3000299, NT2RP3 000397.
       NT2RP3000512, NT2RP3000527, NT2RP3001081, NT2RP3001216, NT2RP3001268, NT2RP 3001355,
       NT2RP3001427, NT2RP3001428, NT2RP3001497, NT2RP3001554, NT2RP3001724, NT2R P3001764,
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       NT2RP3001799, NT2RP3001855, NT2RP3002045, NT2RP3002151, NT2RP3002351, NT2 RP3002352,
       NT2RP3002663, NT2RP3002818, NT2RP3002876, NT2RP3002909, NT2RP3002972, NT 2RP3003061,
       NT2RP3003157, NT2RP3003242, NT2RP3003282, NT2RP3003672, NT2RP3004078, N T2RP3004209,
       NT2RP3004399, NT2RP3004424, NT2RP3004475, NT2RP3004480, NT2RP3004490, NT2RP3004527,
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       NT2RP3004578, NT2RP4000049, NT2RP4000109, NT2RP4000185, NT2RP4000312, NT2RP4000367,
       NT2RP4000398, NT2RP4000457, NT2RP4000855, NT2RP4000879, NT2RP400091 5, NT2RP4000928,
       NT2RP4001095, NT2RP4001150, NT2RP4001213, NT2RP4001483, NT2RP40014 98, NT2RP4001753,
       NT2RP4001838, OVARC1000014, OVARC1000114, OVARC1000139,
       OVARC1000241, OVARC1000440, OVARC1000543, OVARC1000722, OVARC1000771, OVARC1000834,
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       OVARC1000850, OVARC1001113, OVARC1001244, OVARC1001296, OVARC1001357, OVARC100137 2.
       OVARC1001417, OVARC1001496, OVARC1001506, OVARC1001577, OVARC1001668, OVARC10017 13.
       OVARC1001809, OVARC1001880, OVARC1001901, OVARC1001949, OVARC1002050, PLACE1000 133,
       PLACE1000142, PLACE1000184, PLACE1000383, PLACE1000406, PLACE1000420, PLACE100 0583,
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PLACE1000588, PLACE1000706, PLACE1001384, PLACE1001387, PLACE1001517, PLACE10 01602,
       PLACE1001634, PLACE1001771, PLACE1002046, PLACE1002437, PLACE1002474, PLACE1 002665,
       PLACE1002834, PLACE1003302, PLACE1003493, PLACE1003704, PLACE1003888, PLACE 1003903,
       PLACE1003968, PLACE1004197, PLACE1004358, PLACE1004471, PLACE1004681, PLAC E1004751,
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       PLACE1004773, PLACE1005101, PLACE1006040, PLACE1006119, PLACE1006170, PLA CE1006187,
       PLACE1006288, PLACE1006414, PLACE1006438, PLACE1006962, PLACE1007239, PL ACE1007257,
       PLACE1007488, PLACE1007511, PLACE1007737, PLACE1008000, PLACE1008402, P LACE1008643,
       PLACE1008696, PLACE1008941, PLACE1009027, PLACE1009200, PLACE1009230, PLACE1009298.
       PLACE1009319, PLACE1009444, PLACE1009524, PLACE1010083, PLACE1010089, PLACE1010599,
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       PLACE1010833, PLACE1010856, PLACE1011054, PLACE1011057, PLACE101116 5, PLACE1011229,
       PLACE1011263, PLACE1011371, PLACE2000072, PLACE2000399, PLACE20004 38, PLACE2000458,
       PLACE3000004, PLACE3000242, PLACE3000352, PLACE3000353,
       PLACE3000455, PLACE3000475, PLACE4000009, PLACE4000014, PLACE4000052, PLACE4000320,
       PLACE4000581, SKNMC1000050, THYRO1000026, THYRO1000173, THYRO1000186, THYRO100032 0,
       THYRO1000327, THYRO1000387, THYRO1000471, THYRO1000934, THYRO1001458, THYRO10015 95,
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       THYRO1001605, THYRO1001617, THYRO1001671, Y79AA1000037, Y79AA1000469, Y79AA1000 539,
       Y79AA1000560, Y79AA1000734, Y79AA1000985, Y79AA1001105, Y79AA1001391, Y79AA100 1394,
       Y79AA1001548, Y79AA1001613, Y79AA1001874, Y79AA1002258, Y79AA1002416, Y79AA10 02482.
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       (6) The following 301 clones were predicted to encode a protein in the category of enzyme or proteins associated
       with metabolism (including the clones belonging to plural categories).
       HEMBA1000012, HEMBA1000459, HEMBA1000852, HEMBA1001059, HEMBA1001257, HEMBA1001620,
       HEMBA1001714, HEMBA1001744, HEMBA1001896, HEMBA1002003, HEMBA1002212, HEMBA100225 7,
       HEMBA1002513, HEMBA1002973, HEMBA1003046, HEMBA1003136, HEMBA1003179, HEMBA10032 86,
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       HEMBA1003291, HEMBA1003538, HEMBA1003680, HEMBA1004227, HEMBA1004408, HEMBA1004 734,
       HEMBA1004795, HEMBA1005513, HEMBA1005737, HEMBA1005815, HEMBA1006278, HEMBA100 6347,
       HEMBA1006521, HEMBA1006648, HEMBA1006877, HEMBA1006941, HEMBA1006976, HEMBA10 07121,
       HEMBA1007243, HEMBA1007300, HEMBB1000781, HEMBB1000915, HEMBB1001288, HEMBB1 001429,
       HEMBB1001443, HEMBB1001915, HEMBB1002042, HEMBB1002342, HEMBB1002358, HEMBB 1002635,
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       HEMBB1002683, MAMMA1000020, MAMMA1000085, MAMMA1000672, MAMMA1000841, MAMM A1001008.
       MAMMA1001139, MAMMA1001476, MAMMA1001501, MAMMA1002268, MAMMA1002530, MAM MA1002619.
       MAMMA1002671, MAMMA1002938, NT2RM1000132, NT2RM1000153, NT2RM1000256, NT 2RM1000280.
       NT2RM1000377, NT2RM1000648, NT2RM1000850, NT2RM1000867, NT2RM1000894, N T2RM1001072.
       NT2RM2000013, NT2RM2000124, NT2RM2000191, NT2RM2000368, NT2RM2000371, NT2RM2000577,
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       NT2RM2000594, NT2RM2000599, NT2RM2000951, NT2RM2001238, NT2RM2001664, NT2RM2001700,
       NT2RM2001730, NT2RM2001785, NT2RM2001803, NT2RM2002030, NT2RM400002 4, NT2RM4000155,
       NT2RM4000344, NT2RM4000496, NT2RM4000616, NT2RM4000712, NT2RM40008 95, NT2RM4001313,
       NT2RM4001316, NT2RM4001340, NT2RM4001444, NT2RM4001758.
       NT2RM4001819, NT2RM4001930, NT2RM4002062, NT2RM4002063, NT2RM4002409, NT2RM4002558,
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       NT2RM4002571, NT2RM4002623, NT2RP1000018, NT2RP1000112, NT2RP1000376, NT2RP100060 9,
       NT2RP1000833, NT2RP1000834, NT2RP1001079, NT2RP1001253, NT2RP1001361, NT2RP10015 43,
       NT2RP2000056, NT2RP2000114, NT2RP2000183, NT2RP2000329, NT2RP2000422, NT2RP2000 710,
       NT2RP2000764, NT2RP2000932, NT2RP2001070, NT2RP2001663, NT2RP2001839, NT2RP200 1898,
       NT2RP2002256, NT2RP2002312, NT2RP2002442, NT2RP2002595, NT2RP2002618, NT2RP20 02993,
       NT2RP2003230, NT2RP2003286, NT2RP2003506, NT2RP2003643, NT2RP2003706, NT2RP2 003737,
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       NT2RP2003781, NT2RP2003912, NT2RP2004232, NT2RP2004239, NT2RP2004768, NT2RP 2004791,
       NT2RP2004799, NT2RP2005038, NT2RP2005276, NT2RP2005344, NT2RP2005457, NT2R P2005498,
       NT2RP2005531, NT2RP2005557, NT2RP2005690, NT2RP2005773, NT2RP2005775, NT2 RP2005942,
       NT2RP3000571, NT2RP3000046, NT2RP3000085, NT2RP3000359, NT2RP3000418, NT 2RP3000742,
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       NT2RP3000845, NT2RP3000875, NT2RP3001274, NT2RP3001407, NT2RP3001495, N T2RP3002351,
       NT2RP3002402, NT2RP3002969, NT2RP3003301, NT2RP3003346, NT2RP3003500, NT2RP3003800,
       NT2RP3003825, NT2RP3003831, NT2RP3003914, NT2RP3004155, NT2RP3004209 NT2RP3004669,
       NT2RP3004670, NT2RP4000259, NT2RP4000417, NT2RP4000457, NT2RP400072 4, NT2RP4000855,
       NT2RP4000928, NT2RP4000997, NT2RP4001041, NT2RP4001079, NT2RP40010 95, NT2RP4001219,
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       NT2RP4001345, NT2RP4001375, NT2RP4001483, NT2RP4001592,
       NT2RP4001644, NT2RP4001893, NT2RP4001946, NT2RP4002408, NT2RP5003500, NT2RP5003522,
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OVARC100060, OVARC1000139, OVARC1000309, OVARC1000543, OVARC1000682, OVARC100072 2, OVARC1000885, OVARC1001261, OVARC1001610, OVARC1001713, OVARC1001762, OVARC10018 09,

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OVARC1001942, PLACE1000007, PLACE1000142, PLACE1000420, PLACE1000547, PLACE1000 653, PLACE1001062, PLACE1001603, PLACE1001692, PLACE1001817, PLACE1001989, PLACE100 2991, PLACE1003100, PLACE1003709, PLACE1003885, PLACE1003888, PLACE1003903, PLACE10 03915, PLACE1003968, PLACE1004428, PLACE1004437, PLACE1004751, PLACE1005305, PLACE1 005477, PLACE1005763, PLACE1005804, PLACE1006011, PLACE1016469, PLACE 1006731, PLACE1007729, PLACE1007958, PLACE1008275, PLACE1008356, PLACE1008696, PLAC E1009444, PLACE1010481, PLACE1011046, PLACE10110089, PLACE101096, PLACE101106, PLA CE101152, PLACE1011923, PLACE200021, PLACE2000140, PLACE2000404, PLACE2000411, PLACE300020, PLACE3000156, PLACE3000350, PLACE3000350, PLACE3000353, PLACE4000259, PLACE4000431, PLACE4000558, PLACE4000590, SKNMC1000050, THYRO1000017, THYRO1000197, THYRO1000320, THYRO1000358, THYRO1000471, THYRO1000926, THYRO1000934, THYRO100095 1, THYRO1001406, THYRO1001617, THYRO1001671, Y79AA1001048, Y79AA1001233, Y79AA10015 81, Y79AA1001679, Y79AA1001827, Y79AA1002027, Y79AA1002209, Y79AA1002361, Y79AA1002 416,
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Y79AA1000214, Y79AA1001236,

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- (7) The following 74 clones were predicted to encode a protein in the category of proteins associated with cell division or cell proliferation (including the clones belonging to plural categories).

 HEMBA1001019, HEMBA1001595, HEMBA1002229, HEMBA1002363, HEMBA1003136, HEMBA1003836, HEMBA1004131, HEMBA1005241, HEMBB1000399, HEMBB1001175, HEMBB1001242, MAMMA100182 0, MAMMA1003011, NT2RM1000394, NT2RM2001256, NT2RM4000215, NT2RM4001611, NT2RM40017 14, NT2RP1000733, NT2RP1001177, NT2RP2000077, NT2RP2000346, NT2RP2001397, NT2RP2003 912, NT2RP2005520, NT2RP2005675, NT2RP2005767, NT2RP2005857, NT2RP2006464, NT2RP300 0031, NT2RP3001155, NT2RP3001730, NT2RP3002151, NT2RP3002273, NT2RP3002399, NT2RP30 04507, NT2RP3004578, NT2RP3004594, NT2RP4000210, NT2RP4000498, NT2RP4001207, NT2RP4 001414, NT2RP4001551, OVARC1000066, OVARC1000087, OVARC1000113, OVARC1000846, OVARC 1000876, OVARC1001068, OVARC1001107, OVARC1001113, OVARC1002112, PLACE1000547, PLAC E1001611, PLACE1001691, PLACE1002775, PLACE1002811, PLACE1003709, PLACE1004316, PLA CE1004674, PLACE1004814, PLACE1005287, PLACE1006187, PLACE1006368, PLACE1008808, PL ACE1008947, PLACE1009660, PLACE101720, PLACE1010761, PLACE1010833, PLACE1011056, P LACE3000244,
- (8) The following 92 clones were predicted to encode a protein in the category of proteins associated with cytoskeleton (including the clones belonging to plural categories).
- HEMBA100020, HEMBA1000588, HEMBA1001804, HEMBA1002102, HEMBA1002161, HEMBA1002939, HEMBA1003235, HEMBA1003581, HEMBA1004534, HEMBA1005009, HEMBA1005595, HEMBA100627 2, HEMBA1006737, HEMBA1007018, HEMBB1000404, HEMBB1001282, MAMMA1000173, MAMMA10004 29, MAMMA1001041, MAMMA1001576, MAMMA1001735, MAMMA1002622, MAMMA1002637, MAMMA1003 099, MAMMA1003127, NT2RM1000898, NT2RM1001003, NT2RM1001139, NT2RM2000691, NT2RM400 0030, NT2RM4000167, NT2RM4000689, NT2RM4001741, NT2RM4001776, NT2RM4002109, NT2RP10 00348, NT2RP1000460, NT2RP1000478, NT2RP1001033, NT2RP2000812, NT2RP2001634, NT2RP2 001900, NT2RP2002727, NT2RP2003307, NT2RP2003604, NT2RP2004538, NT2RP2005531, NT2RP 3000868, NT2RP3001216, NT2RP3001272, NT2RP3002688, NT2RP3003061, NT2RP3003138, NT2R P3003230, NT2RP3003282, NT2RP3004454, NT2RP3004569, NT2RP4001126, NT2RP4001927, OVA RC1000437, OVARC1000520, OVARC1000679, OVARC1001731, OVARC1002050, PLACE1002571, PL ACE1002591, PLACE1002655, PLACE1003669, PLACE1004777, PLACE1005287, PLACE1006368, P LACE1007105, PLACE1010310, PLACE1010743, PLACE1010960, PLACE1011263, PLACE2000039, PLACE2000216, PLACE2000274, PLACE3000145, PLACE3000322, PLACE4000009, PLACE4000612, THYRO1000072, THYRO1000666, THYRO1001121, THYRO1001458, Y79AA1000368, Y79AA100083 3, Y79AA1000962, Y79AA1000985, Y79AA1002208,

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(9) The following 280 clones were predicted to encode a protein in the category of proteins associated with RNA synthesis (including the clones belonging to plural categories).

HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000591, HEMBA1000851, HEMBA1001137, HEMBA1001510, HEMBA1001579, HEMBA1001800, HEMBA1001809, HEMBA100181 9, HEMBA1001847, HEMBA1002035, HEMBA1002092, HEMBA1002569, HEMBA1002609, HEMBA10034 18, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA1003953, HEMBA1004 097, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA1004669, HEMBA1004758, HEMBA100 5359, HEMBA1005528, HEMBA1005548, HEMBA1005931, HEMBA1006253, HEMBA1006278, HEMBA10 06359,

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HEMBA1006559, HEMBA1006639, HEMBB1000250, HEMBB1000789, HEMBB1001011, HEMBB1 001314.
        HEMBB1001482, HEMBB1001908, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAMMA 1000183,
       MAMMA1000388, MAMMA1001059, MAMMA1001633, MAMMA1001820, MAMMA1001837, MAMM A1002650.
       MAMMA1002937, NT2RM1000055, NT2RM1000187, NT2RM1000666, NT2RM1000691, NT2 RM1000852,
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       NT2RM1000894, NT2RM1001092, NT2RM2000013, NT2RM2000735, NT2RM2001035, NT 2RM2001424,
       NT2RM2001575, NT2RM2001605, NT2RM2001670, NT2RM2001771, NT2RM2001797, N T2RM2001989,
       NT2RM2002100, NT2RM2002580, NT2RM4000024, NT2RM4000104, NT2RM4000191, NT2RM4000421,
       NT2RM4000471, NT2RM4000496, NT2RM4000531, NT2RM4000734, NT2RM4000751, NT2RM4000996.
       NT2RM4001140, NT2RM4001200, NT2RM4001483, NT2RM4001592, NT2RM400174 6, NT2RM4001783,
       NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM4001979, NT2RM40020 93, NT2RM4002479,
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       NT2RP1000111, NT2RP1000272, NT2RP1000333, NT2RP1000470.
       NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1001080, NT2RP2000008, NT2RP2000133,
       NT2RP2000161, NT2RP2000297, NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP200144 9,
       NT2RP2002099, NT2RP2002252, NT2RP2002503, NT2RP2002591, NT2RP2002928, NT2RP20031 25,
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       NT2RP2003228, NT2RP2003293, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2RP2004 187,
       NT2RP2004568, NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005126, NT2RP200 5168,
       NT2RP2005239, NT2RP2005287, NT2RP2005496, NT2RP2005539, NT2RP2005722, NT2RP20 05763,
       NT2RP2005942, NT2RP2006238, NT2RP2006464, NT2RP3000050, NT2RP3000361, NT2RP3 000397,
       NT2RP3000512, NT2RP3000527, NT2RP3000590, NT2RP3000632, NT2RP3000917, NT2RP 3001057,
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       NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001398, NT2RP3001399, NT2R P3001671,
       NT2RP3001672, NT2RP3001688, NT2RP3001712, NT2RP3001792, NT2RP3001855, NT2 RP3002165,
       NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193, NT2RP3003251, NT 2RP3003327,
       NT2RP3004013, NT2RP3004258, NT2RP3004504, NT2RP3004566, NT2RP3004572, N T2RP3004594,
       NT2RP3004617, NT2RP4000078, NT2RP4000111, NT2RP4000210, NT2RP4000398, NT2RP4000449,
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       NT2RP4000518, NT2RP4000614, NT2RP4000787, NT2RP4000865, NT2RP4000997 , NT2RP4001029,
       NT2RP4001080, NT2RP4001148, NT2RP4001213, NT2RP4001433, NT2RP400152 9, NT2RP4001551,
       NT2RP4001638, NT2RP4001753, NT2RP4001790, NT2RP4001938, NT2RP40020 78, OVARC1000479,
       OVARC1000846, OVARC1001381, OVARC1001555, OVARC1001577,
        OVARC1001949, PLACE1000066, PLACE1000133, PLACE1000583, PLACE1000596, PLACE1000706,
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       PLACE1001118, PLACE1001238, PLACE1001304, PLACE1001602, PLACE1001632, PLACE100245 0,
       PLACE1002532, PLACE1002834, PLACE1003190, PLACE1003302, PLACE1003605, PLACE10037 04,
       PLACE1003738, PLACE1003885, PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004 681,
       PLACE1004902, PLACE1005646, PLACE1005876, PLACE1006196, PLACE1006414, PLACE100 6438,
       PLACE1006482, PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10 07955,
35
       PLACE1007969, PLACE1008627, PLACE1008941, PLACE1009099, PLACE1009246, PLACE1 009398,
       PLACE1010053, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010580, PLACE 1010702,
       PLACE1010771, PLACE1010870, PLACE1011576, PLACE2000072, PLACE2000359, PLAC E4000128,
       PLACE4000156, PLACE4000259, PLACE4000431, THYRO1000121, THYRO1000242, THY RO1000501,
        THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000349, Y7 9AA1000539,
        Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1001105, Y79AA1001299, Y 79AA1001391,
40
        Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA1001866, Y79AA1001963, Y79AA1002093,
        Y79AA1002472, Y79AA1002482,
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(10) The following 352 clones were predicted to encode a protein in the category of nuclear protein (including the clones belonging to plural categories).

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HEMBA100005, HEMBA1000201, HEMBA1000216, HEMBA1000488, HEMBA1000561, HEMBA1000827, HEMBA1000851, HEMBA1001088, HEMBA1001137, HEMBA1001476, HEMBA1001510, HEMBA100157 9, HEMBA1001800, HEMBA1001809, HEMBA1001819, HEMBA1001847, HEMBA1002035, HEMBA10020 92, HEMBA1002363, HEMBA1002495, HEMBA1002569, HEMBA1002609, HEMBA1002999, HEMBA1003 148, HEMBA1003418, HEMBA1003545, HEMBA1003568, HEMBA1003684, HEMBA1003760, HEMBA100 3783, HEMBA1003953, HEMBA100497, HEMBA1004321, HEMBA1004353, HEMBA1004479, HEMBA10 04669, HEMBA1004758, HEMBA1005359, HEMBA1005475, HEMBA1005528, HEMBA1005548, HEMBA1 005931, HEMBA1006253, HEMBA1006278, HEMBA1006359, HEMBA1006474, HEMBA1006559, HEMBA 1006914, HEMBB1001250, HEMBB1000399, HEMBB100789, HEMBB1001011, HEMBB1001314, HEMB B1001482, HEMBB1001908, HEMBB1001915, HEMBB1002134, HEMBB1002217, MAMMA1000155, MAM MA1000183, MAMMA1000388, MAMMA1000731, MAMMA1001633, MAMMA1001820, MAMMA1001837, NA MMA100257, NT2RM1000394, NT2RM1000666, NT2RM1000691, NT2RM1000852, NT2RM1000894, NT2RM1001092,

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NT2RM2000013, NT2RM2000735, NT2RM2001035, NT2RM2001575, NT2RM2001605, NT2RM2001635,
       NT2RM2001670, NT2RM2001771, NT2RM2001797, NT2RM2001823, NT2RM200193 6, NT2RM2001989,
       NT2RM2002100. NT2RM2002580, NT2RM4000024, NT2RM4000085, NT2RM40001 04, NT2RM4000191.
       NT2RM4000215, NT2RM4000290, NT2RM4000421, NT2RM4000496,
       NT2RM4000531, NT2RM4000734, NT2RM4000751, NT2RM4000996, NT2RM4001140, NT2RM4001200.
       NT2RM4001217, NT2RM4001382, NT2RM4001483, NT2RM4001592, NT2RM4001605, NT2RM400161 1,
       NT2RM4001746, NT2RM4001783, NT2RM4001823, NT2RM4001828, NT2RM4001858, NT2RM40019 79,
       NT2RM4002093, NT2RM4002146, NT2RM4002527, NT2RP1000035, NT2RP1000111, NT2RP1000 333.
       NT2RP1000493, NT2RP1000574, NT2RP1000721, NT2RP1000860, NT2RP1000966, NT2RP100 1177,
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       NT2RP2000008, NT2RP2000126, NT2RP2000133, NT2RP2000161, NT2RP2000297, NT2RP20 00414.
       NT2RP2000931, NT2RP2001174, NT2RP2001233, NT2RP2001420, NT2RP2001449, NT2RP2 001536.
       NT2RP2001762, NT2RP2002185, NT2RP2002252, NT2RP2002259, NT2RP2002503, NT2RP 2002504,
       NT2RP2002591, NT2RP2002993, NT2RP2003125, NT2RP2003228, NT2RP2003277, NT2R P2003286,
       NT2RP2003293, NT2RP2003308, NT2RP2003564, NT2RP2003714, NT2RP2004013, NT2 RP2004187,
       NT2RP2004961, NT2RP2005003, NT2RP2005037, NT2RP2005287, NT2RP2005496, NT 2RP2005520,
       NT2RP2005539, NT2RP2005722, NT2RP2005767, NT2RP2005857, NT2RP2005942, N T2RP2006238,
       NT2RP2006464, NT2RP3000031, NT2RP3000050, NT2RP3000361, NT2RP3000397, NT2RP3000512,
       NT2RP3000527, NT2RP1000590, NT2RP3000603, NT2RP3000632, NT2RP3000917, NT2RP3001057,
       NT2RP3001120, NT2RP3001155, NT2RP3001268, NT2RP3001274, NT2RP300139 8, NT2RP3001399,
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       NT2RP3001427, NT2RP3001428, NT2RP3001587, NT2RP3001671, NT2RP30016 72, NT2RP3001688,
       NT2RP3001712, NT2RP3001724, NT2RP3001792, NT2RP3001855,
       NT2RP3002056, NT2RP3002165, NT2RP3002377, NT2RP3002399, NT2RP3003157, NT2RP3003193,
       NT2RP3003212, NT2RP3003251, NT2RP3003327, NT2RP3004206, NT2RP3004209, NT2RP300425 8,
       NT2RP3004566, NT2RP3004572, NT2RP3004594, NT2RP3004617, NT2RP4000078, NT2RP40001 11.
       NT2RP4000150, NT2RP4000210, NT2RP4000398, NT2RP4000449, NT2RP4000457, NT2RP4000 518,
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       NT2RP4000787, NT2RP4000865, NT2RP4000997, NT2RP4001029, NT2RP4001080, NT2RP400 1148,
       NT2RP4001207, NT2RP4001213, NT2RP4001433, NT2RP4001529, NT2RP4001551, NT2RP40 01638,
       NT2RP4001753, NT2RP4001790. NT2RP4001938, NT2RP4002078, OVARC1000006, OVARC1 000087,
       OVARC1000139, OVARC1000326, OVARC1000440, OVARC1000479, OVARC1000846, OVARC 1000883,
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       OVARC1001180, OVARC1001244, OVARC1001381, OVARC1001555, OVARC1001577, OVAR C1001702,
       OVARC1001949, OVARC1002112, PLACE1000066, PLACE1000133, PLACE1000406, PLA CE1000583,
       PLACE1000596, PLACE1000706, PLACE1001118, PLACE1001238, PLACE1001304, PL ACE1001602,
       PLACE1001611, PLACE1001632, PLACE1002450, PLACE1002532, PLACE1002834, P LACE1003100,
       PLACE1003190, PLACE1003302, PLACE1003605, PLACE1003704, PLACE1003738, PLACE1003885,
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       PLACE1004471, PLACE1004510, PLACE1004564, PLACE1004681, PLACE1004902, PLACE1005287,
       PLACE1005876, PLACE1006011, PLACE1006119, PLACE1006414, PLACE100643 8, PLACE1006482,
       PLACE1007239, PLACE1007346, PLACE1007544, PLACE1007598, PLACE10079 55, PLACE1007969,
       PLACE1008044, PLACE1008177, PLACE1008603, PLACE1008627,
       PLACE1008941, PLACE1008947, PLACE1009099, PLACE1009113, PLACE1009246, PLACE1009398.
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       PLACE1010152, PLACE1010194, PLACE1010261, PLACE1010491, PLACE1010702, PLACE101072 0,
       PLACE1010771, PLACE1010870, PLACE1011056, PLACE1011229, PLACE1011576, PLACE10116 64,
       PLACE2000072, PLACE2000359, PLACE2000411, PLACE4000014, PLACE4000128, PLACE4000 156,
       PLACE4000259, PLACE4000261, PLACE4000431, THYRO1000121, THYRO1000242, THYRO100 0501,
       THYRO1000585, THYRO1001100, THYRO1001189, Y79AA1000037, Y79AA1000214, Y79AA10 00539,
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       Y79AA1000589, Y79AA1000627, Y79AA1000705, Y79AA1000752, Y79AA1000784, Y79AA1 001105,
       Y79AA1001299, Y79AA1001391, Y79AA1001533, Y79AA1001613, Y79AA1001848, Y79AA 1001866,
       Y79AA1001963, Y79AA1002093, Y79AA1002472, Y79AA1002482,
       (11) The following 111 clones were predicted to encode a protein in the category of proteins associated with protein
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       synthesis or transport (including the clones belonging to plural categories).
       HEMBA1000012, HEMBA1000542, HEMBA1001913, HEMBA1003773, HEMBA1004202, HEMBA1004356,
       HEMBA1004847, HEMBA1005047, HEMBA1005202, HEMBA1005963, HEMBA1006310, HEMBA100665 2,
       HEMBA1006914, HEMBB1000217, HEMBB1000725, HEMBB1001346, HEMBB1001831, MAMMA10000 85,
       MAMMA1000734, MAMMA1002170, MAMMA1002236, MAMMA1002598, NT2RM1000661, NT2RM2000 504,
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       NT2RM2000577, NT2RM2001131, NT2RM4000155, NT2RM4001444, NT2RM4002062, NT2RM400 2205,
       NT2RM4002623, NT2RP1000326, NT2RP1000947, NT2RP1001569, NT2RP2000710, NT2RP20 00880,
       NT2RP2001290, NT2RP2001511, NT2RP2001660, NT2RP2002185, NT2RP2002606, NT2RP2 002959,
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NT2RP2002980, NT2RP2003158, NT2RP2003760, NT2RP2004194, NT2RP2004791, NT2RP 2005012,

NT2RP2005116, NT2RP2005204, NT2RP2006565, NT2RP2006598, NT2RP3000047, NT2R P3000366, NT2RP3000759, NT2RP3000968, NT2RP3001587, NT2RP3002529, NT2RP3003385, NT2 RP3003589, NT2RP3003876, NT2RP3004618, NT2RP4000370, NT2RP4000524, NT2RP4000528, NT 2RP4001041, NT2RP4001574, NT2RP4001592, NT2RP4002047, OVARC1000004, OVARC1000771, O VARC1000862, OVARC1001180, OVARC1001342, OVARC1001766, OVARC1002138, PLACE1000061, PLACE1000492, PLACE1000863, PLACE1001748, PLACE1002090, PLACE1003394, PLACE1003915, PLACE1004149, PLACE1004743, PLACE1005557, PLACE1005813, PLACE1006488, PLACE100653 1, PLACE1006615, PLACE1007706, PLACE1008273, PLACE1008790, PLACE1008813, PLACE10097 21, PLACE1009763, PLACE1009845, PLACE1010074, PLACE1011109, PLACE1011477,

PLACE2000404, PLACE3000059, PLACE3000121, PLACE4000558, PLACE4000654, SKNMC1000011, THYRO1000288, THYRO1000974, Y79AA1000346, Y79AA1000968, Y79AA1002209.

(12) The following 23 clones were predicted to encode a protein in the category of proteins associated with cellular defense (including the clones belonging to plural categories).

HEMBA100005, HEMBA1000531, HEMBB1000217, HEMBB1000399, MAMMA1000734, NT2RP1000493, NT2RP1000493, NT2RP200006, NT2RP2000045, NT2RP2001536, NT2RP2005204, NT2RP300059 0, NT2RP3001426, NT2RP3002056, NT2RP3004262, NT2RP4001638, OVARC1001900, PLACE10056 11, PLACE1006958, PLACE1008275, PLACE1009113, PLACE4000014, Y79AA1002139,

(13) The following 23 clones were predicted to encode a protein in the category of proteins associated with development or growth (including the clones belonging to plural categories).
 HEMBB1001802, HEMBB1002442, NT2RM2002142, NT2RM4001047, NT2RP2003308, NT2RP2004816, NT2RP3000994, NT2RP3001340, NT2RP3004206, NT2RP3004472, NT2RP4000246, NT2RP400100 6, OVARC1000304, OVARC1000746, OVARC1000996, PLACE1006037, PLACE1009622, PLACE10116 64, Y79AA1001692.

Table 9
Selected clones having the maximal ATGpr1 score of 0.3 or higher (3690 clones).

clone	name of	maximal
name	sequence	ATGpr1
		score
HEMBA1000005	F-HEMBA1000005	0. 84
HEMBA1000012	F-HEMBA1000012	0. 56
HEMBA1000020	F-HEMBA1000020	0.94
HEMBA1000030	F-HEMBA1000030	0.44
HEMBA1000046	F-HEMBA1000046	0.50
HEMBA1000050	F-HEMBA1000050	0.94
HEMBA1000076	F-HEMBA1000076	0.48
HEMBA1000129	F-HEMBA1000129	0.74
HEMBA1000141	F-HEMBA1000141	0.55
HEMBA1000150	F-HEMBA1000150	0. 72
HEMBA1000156	F-HEMBA1000156	0.94
HEMBA1000158	F-HEMBA1000158	0. 62
HEMBA1000168	F-HEMBA1000168	0.94
HEMBA1000185	F-HEMBA1000185	0.86
HEMBA1000193	F-HEMBA1000193	0.94

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	HEMBA1000201	F-HEMBA1000201	0.83
	HEMBA1000216	F-HEMBA1000216	0.32
	HEMBA1000227	F-HEMBA1000227	0.72
5	HEMBA1000231	F-HEMBA1000231	0, 34
	HEMBA1000244	F-HEMBA1000244	0.51
	HEMBA1000288	F-HEMBA1000288	0.36
	HEMBA1000290	F-HEMBA1000290	0. 45
10	HEMBA1000302	F-HEMBA1000302	0.71
	HEMBA1000303	F-HEMBA1000303	0. 42
	HEMBA1000304	F-HEMBA1000304	0. 45
	HEMBA1000307	F-HEMBA1000307	0. 48
15	HEMBA1000327	F-HEMBA1000327	0. 94
	HEMBA1000356	F-HEMBA1000356	0.90
	HEMBA1000369	F-HEMBA1000369,	0. 94
	HEMBA1000387	F-HEMBA1000387	0.66
20	HEMBA1000392	F-HEMBA1000392	0.60
-	HEMBA1000396	F-HEMBA1000396	0.81
	HEMBA1000456	F-HEMBA1000456	0.90
•	HEMBA1000460	F-HEMBA1000460	0.31
25	HEMBA1000488	F-HEMBA1000488	0.47
	HEMBA1000490	F-HEMBA1000490	0.89
	HEMBA1000491	F-HEMBA1000491	0.94
	HEMBA1000501	F-HEMBA1000501	0.67
30	HEMBA1000505	F-HEMBA1000505	0. 65
30	HEMBA1000508	F-HEMBA1000508	0. 39
	HEMBA1000520	F-HEMBA1000520	0. 94
	HEMBA1000523	F-HEMBA1000523	0. 86
25	HEMBA1000531	F-HEMBA1000531	0. 49
35	HEMBA1000534	F-HEMBA1000534	0.52
	HEMBA1000542	F-HEMBA1000542	0.81
	HEMBA1000555	F-HEMBA1000555	0.67
40	HEMBA1000561	F-HEMBA1000561	0.34
40	HEMBA1000568	F-HEMBA1000568 F-HEMBA1000588	0.66
	HEMBA1000588 HEMBA1000591	F-HEMBA1000591	0.79
	HEMBA1000591	F-HEMBA1000592	0. 92 0. 73
	HEMBA1000594	F-HEMBA1000594	0.73
45	HEMBA1000594	F-HEMBA1000608	0. 94
	HEMBA1000636	F-HEMBA1000636	0. 94
	HEMBA1000637	F-HEMBA1000637	0. 92
	HEMBA1000657	F-HEMBA1000657	0. 32
50	HEMBA1000682	F-HEMBA1000682	0. 94
	HEMBA1000686	F-HEMBA1000686	0.90
	HEMBA1000719	F-HEMBA1000719	0. 94
	HEMBA1000727	F-HEMBA1000727	0.40
55	HEMBA1000752	F-HEMBA1000752	0. 31

	HEMBA1000817	F-HEMBA1000817	0.94
	HEMBA1000851	F-HEMBA1000851	0. 63
	HEMBA1000867	F-HEMBA1000867	0.68
5	HEMBA1000869	F-HEMBA1000869	0.82
	HEMBA1000872	F-HEMBA1000872	0.54
	HEMBA1000910	F-HEMBA1000910	0.91
	HEMBA1000918	F-HEMBA1000918	0.38
10	HEMBA1000919	F-HEMBA1000919	0.83
	HEMBA1000942	F-HEMBA1000942	0. 31
	HEMBA1000946	F-HEMBA1000946	0. 75
	HEMBA1000968	F-HEMBA1000968	0.39
15	HEMBA1000971	F-HEMBA1000971	0.74
	HEMBA1000975	F-HEMBA1000975	0.46
	HEMBA1000986	F-HEMBA1000986,	0.89
	HEMBA1001008	F-HEMBA1001008	0. 47
20	HEMBA1001009	F-HEMBA1001009	0.86
	HEMBA1001022	F-HEMBA1001022	0.57
	HEMBA1001043	F-HEMBA1001043	0.79
	HEMBA1001052	F-HEMBA1001052	0.83
25	HEMBA1001059	F-HEMBA1001059	0.78
	HEMBA1001080	F-HEMBA1001080	0.77
	HEMBA1001085	F-HEMBA1001085	0.92
	HEMBA1001088	F-HEMBA1001088	0.94
30	HEMBA1001109	F-HEMBA1001109	0.50
	HEMBA1001122	F-HEMBA1001122	0. 74
	HEMBA1001133	F-HEMBA1001133	0. 45
	HEMBA1001137	F-HEMBA1001137	0.37
35	HEMBA1001140	F-HEMBA1001140	0. 43
30	HEMBA1001174	F-HEMBA1001174	0. 44
	HEMBA1001197	F-HEMBA1001197	0.65
	HEMBA1001213	F-HEMBA1001213	0. 61
40	HEMBA1001235	F-HEMBA1001235	0. 94
40	HEMBA1001247	F-HEMBA1001247	0.31
	HEMBA1001257	F-HEMBA1001257	0.92
	HEMBA1001281 HEMBA1001286	F-HEMBA1001281 F-HEMBA1001286	0.94
	HEMBA1001289	F-HEMBA1001289	0. 49 0. 94
45	HEMBA1001302	F-HEMBA1001289	0. 62
	HEMBA1001303	F-HEMBA1001302	0. 62
	HEMBA1001310	F-HEMBA1001310	0. 79
	HEMBA1001316	F-HEMBA1001326	0. 59
50	HEMBA1001351	F-HEMBA1001351	0. 65
	HEMBA1001361	F-HEMBA1001361	0. 94
	HEMBA1001377	F-HEMBA1001377	0. 87
	HEMBA1001387	F-HEMBA1001387	0. 94
55	HEMBA1001388	F-HEMBA1001388	0.57
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	HEMBA1001398	F-HEMBA1001398	0. 74
	HEMBA1001405	F-HEMBA1001405	0. 61
	HEMBA1001407	F-HEMBA1001407	0. 43
5	HEMBA1001413	F-HEMBA1001413	0.40
	HEMBA1001415	F-HEMBA1001415	0.94
	HEMBA1001446	F-HEMBA1001446	0.90
	HEMBA1001450	F-HEMBA1001450	0.40
10	HEMBA1001455	F-HEMBA1001455	0. 73
	HEMBA1001476	F-HEMBA1001476	0.65
	HEMBA1001497	F-HEMBA1001497	0. 48
	HEMBA1001510	F-HEMBA1001510	0.36
15	HEMBA1001526	F-HEMBA1001526	0.66
	HEMBA1001533	F-HEMBA1001533	0, 46
	HEMBA1001569	F-HEMBA1001569,	
	HEMBA1001570	F-HEMBA1001570	0. 53
20	HEMBA1001579	F-HEMBA1001579	0. 82
	HEMBA1001581	F-HEMBA1001581	0. 80
	HEMBA1001595	F-HEMBA1001595	0. 72
	HEMBA1001620	F-HEMBA1001620	0.94
25	HEMBA1001635	F-HEMBA1001635	0. 87
20	HEMBA1001640	F-HEMBA1001640	0.82
	HEMBA1001647	F-HEMBA1001647	0. 91
	HEMBA1001655	F-HEMBA1001655	0.39
•	HEMBA1001661	F-HEMBA1001661	0.86
30	HEMBA1001672	F-HEMBA1001672	0. 94
	HEMBA1001702	F-HEMBA1001702	0. 72
	HEMBA1001711	F-HEMBA1001711	0.74
	HEMBA1001714	F-HEMBA1001714	0.89
35	HEMBA1001723	F-HEMBA1001723	0. 94
	HEMBA1001731	F-HEMBA1001731	0.37
	HEMBA1001744	F-HEMBA1001744	0.31
	HEMBA1001746	F-HEMBA1001746	0. 39
40	HEMBA1001781	F-HEMBA1001781	0.67
	HEMBA1001800	F-HEMBA1001800	0.69
	HEMBA1001804	F-HEMBA1001804	0. 85
	HEMBA1001809	F-HEMBA1001809	0. 78
45	HEMBA1001815	F-HEMBA1001815	0. 49
	HEMBA1001819	F-HEMBA1001819	0. 90
	HEMBA1001822	F-HEMBA1001822	0. 80
	HEMBA1001824	F-HEMBA1001824	0. 68
50	HEMBA1001847	F-HEMBA1001847	0. 46
	HEMBA1001864	F-HEMBA1001864	0.34
	HEMBA1001866	F-HEMBA1001866	0. 53
	HEMBA1001869	F-HEMBA1001869	0. 69
55	HEMBA1001896	F-HEMBA1001896	0. 48
	HEMBA1001910	F-HEMBA1001910	0. 84

	HEMBA1001912	F-HEMBA1001912	0. 48
	HEMBA1001913	F-HEMBA1001913	0.94
	HEMBA1001915	F-HEMBA1001915	0.41
5	HEMBA1001921	F-HEMBA1001921	0. 94
	HEMBA1001939	F-HEMBA1001939	0. 87
	HEMBA1001950	F-HEMBA1001950	0. 36
	HEMBA1001967	F-HEMBA1001967	0. 94
10	HEMBA1001987	F-HEMBA1001987	0.41
	HEMBA1002018	F-HEMBA1002018	0. 41
	HEMBA1002035	F-HEMBA1002035	0. 36
	HEMBA1002049	F-HEMBA1002049	0. 39
15	HEMBA1002084	F-HEMBA1002084	0. 45
•	HEMBA1002092	F-HEMBA1002092	0. 94
	HEMBA1002102	F-HEMBA1002102,	
	HEMBA1002102	F-HEMBA1002119	0. 60
	HEMBA1002115	F-HEMBA1002125	0. 88
20	HEMBA1002150	F-HEMBA1002125	0. 31
	HEMBA1002151	F-HEMBA1002151	0. 82
	HEMBA1002161	F-HEMBA1002161	0. 94
	HEMBA1002177	F-HEMBA1002177	0. 34
25	HEMBA1002177	F-HEMBA1002177	0. 43
	HEMBA1002191	F-HEMBA1002191	0. 43
	HEMBA1002199	F-HEMBA1002199	0. 85
	HEMBA1002199	F-HEMBA1002199	0. 39
30	HEMBA1002212	F-HEMBA1002215	0. 94
	HEMBA1002229	F-HEMBA1002219	0. 83
		F-HEMBA1002237	0. 86
	HEMBA1002237 HEMBA1002241	F-HEMBA1002241	0. 66
35	HEMBA1002241	F-HEMBA1002241	0. 38
	HEMBA1002267	F-HEMBA1002267	0. 33
	HEMBA1002341	F-HEMBA1002341	0. 53
	HEMBA1002341	F-HEMBA1002341	0. 88
40	HEMBA1002417	F-HEMBA1002417	0. 83
•	HEMBA1002419	F-HEMBA1002419	0. 92
	HEMBA1002439	F-HEMBA1002430	0. 45
	HEMBA1002439	F-HEMBA1002439	0. 43
	HEMBA1002458	F-HEMBA1002458	0. 63
45	HEMBA1002460	F-HEMBA1002460	0.47
	HEMBA1002462	F-HEMBA1002462	0. 55
	HEMBA1002469	F-HEMBA1002469	0. 79
	HEMBA1002475	F-HEMBA1002475	0. 79
50		F-HEMBA1002477	0. 45
	HEMBA1002477	F-HEMBA1002477	0. 45
	HEMBA1002495	,	
	HEMBA1002503	F-HEMBA1002503	0. 94
55	HEMBA1002508		0. 33
	HEMBA1002513	F-HEMBA1002513	0. 41

	HEMBA1002515	F-HEMBA1002515	0. 94
	HEMBA1002542	F-HEMBA1002542	0. 46
	HEMBA1002547	F-HEMBA1002547	0. 45
5			
	HEMBA1002569	F-HEMBA1002569	
	HEMBA1002583	F-HEMBA1002583	0.86
	HEMBA1002609	F-HEMBA1002609	0. 75
10	HEMBA1002624	F-HEMBA1002624	0. 59
,,,	HEMBA1002688	F-HEMBA1002688	0.88
	HEMBA1002696	F-HEMBA1002696	0. 82
	HEMBA1002703	F-HEMBA1002703	0.94
	HEMBA1002746	F-HEMBA1002746	0. 33
15	HEMBA1002750	F-HEMBA1002750	0. 53
	HEMBA1002768	F-HEMBA1002768	0.94
	HEMBA1002770	F-HEMBA1002770,	
	HEMBA1002777	F-HEMBA1002777	
20	HEMBA1002779	F-HEMBA1002779	0. 92
	HEMBA1002794	F-HEMBA1002794	0.86
	HEMBA1002810	F-HEMBA1002810	0. 87
	HEMBA1002816	F-HEMBA1002816	0. 91
25	HEMBA1002818	F-HEMBA1002818	0.94
	HEMBA1002850	F-HEMBA1002850	0.57
	HEMBA1002863	F-HEMBA1002863	0. 48
	HEMBA1002876	F-HEMBA1002876	0.86
30	HEMBA1002935	F-HEMBA1002935	0.91
	HEMBA1002937	F-HEMBA1002937	0.61
	HEMBA1002939	F-HEMBA1002939	0. 32
	HEMBA1002951	F-HEMBA1002951	0.83
35	HEMBA1002954	F-HEMBA1002954	0. 67
33	HEMBA1002970	F-HEMBA1002970	0. 33
	HEMBA1002971	F-HEMBA1002971	0. 57
	HEMBA1002973	F-HEMBA1002973	0. 33
	HEMBA1002997	F-HEMBA1002997	0. 46
40	HEMBA1002999	F-HEMBA1002999	0. 49
	HEMBA1003021	F-HEMBA1003021	0. 74
		F-HEMBA1003033	0. 94
	HEMBA1003035	F-HEMBA1003035	0. 61
45	HEMBA1003041	F-HEMBA1003041	0.40
	HEMBA1003046	F-HEMBA1003046	0. 94
	HEMBA1003067	F-HEMBA1003067	0. 34
	HEMBA1003077	F-HEMBA1003077	0. 83
50	HEMBA1003078	F-HEMBA1003078	0.89
	HEMBA1003079	F-HEMBA1003079	0. 49
	HEMBA1003096	F-HEMBA1003096	0. 34
	HEMBA1003117	F-HEMBA1003117	0. 94
55	HEMBA1003129	F-HEMBA1003129	0. 81
	HEMBA1003136	F-HEMBA1003136	0. 77

	HEMBA1003148	F-HEMBA1003148	0. 91
	HEMBA1003175	F-HEMBA1003175	0.41
_	HEMBA1003179	F-HEMBA1003179	0. 42
5	HEMBA1003199	F-HEMBA1003199	0.84
	HEMBA1003222	F-HEMBA1003222	0. 47
	HEMBA1003235	F-HEMBA1003235	0.94
	HEMBA1003250	F-HEMBA1003250	0.39
10	HEMBA1003257	F-HEMBA1003257	0.88
	HEMBA1003281	F-HEMBA1003281	0. 67
	HEMBA1003286	F-HEMBA1003286	0. 57
	HEMBA1003291	F-HEMBA1003291	0.89
15	HEMBA1003322	F-HEMBA1003322	0.60
	HEMBA1003327	F-HEMBA1003327	0.88
	HEMBA1003369	F-HEMBA1003369,	0.89
	HEMBA1003370	F-HEMBA1003370	0. 75
20	HEMBA1003380	F-HEMBA1003380	0.68
	HEMBA1003395	F-HEMBA1003395	0.88
	HEMBA1003402	F-HEMBA1003402	0, 45
	HEMBA1003403	F-HEMBA1003403	0. 44
25	HEMBA1003408	F-HEMBA1003408	0.34
	HEMBA1003417	F-HEMBA1003417	0. 78
	HEMBA1003418	F-HEMBA1003418	0, 94
	HEMBA1003433	F-HEMBA1003433	0. 70
30	HEMBA1003447	F-HEMBA1003447	0. 49
	HEMBA1003461	F-HEMBA1003461	0. 85
	HEMBA1003463	F-HEMBA1003463	0. 59
	HEMBA1003480	F-HEMBA1003480	0. 94
35	HEMBA1003528	F-HEMBA1003528	0. 90
	HEMBA1003538	F-HEMBA1003538	0. 47
	HEMBA1003545	F-HEMBA1003545	0. 74
	HEMBA1003555	F-HEMBA1003555	0. 94
40	HEMBA1003556	F-HEMBA1003556	0. 44
	HEMBA1003560	F-HEMBA1003560	0. 40
	HEMBA1003568	F-HEMBA1003568	0. 94
	HEMBA1003569	F-HEMBA1003569	0. 82
45	HEMBA1003581 HEMBA1003591	F-HEMBA1003581	0. 83
43	HEMBA1003591	F-HEMBA1003591 F-HEMBA1003615	0. 68
	HEMBA1003617	F-HEMBA1003617	0.94
	HEMBA1003621	F-HEMBA1003621	0.88
50	HEMBA1003645	F-HEMBA1003645	0. 47 0. 75
50	HEMBA1003646	F-HEMBA1003646	0. 73
	HEMBA1003662	F-HEMBA1003662	0. 61
	HEMBA1003667	F-HEMBA1003667	0. 54
	HEMBA1003679	F-HEMBA1003679	0. 72
55	HEMBA1003680	F-HEMBA1003680	0. 72
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	HEMBA1003684	F-HEMBA1003684	0. 94
	HEMBA1003690	F-HEMBA1003690	0. 93
	HEMBA1003692	F-HEMBA1003692	0. 46
5	HEMBA1003711	F-HEMBA1003711_	0. 94
	HEMBA1003720	F-HEMBA1003720	0. 94
	HEMBA1003729	F-HEMBA1003729	0. 73
	HEMBA1003742	F-HEMBA1003742	0. 94
10	HEMBA1003760	F-HEMBA1003760	0. 75
	HEMBA1003783	F-HEMBA1003783	0. 31
	HEMBA1003799	F-HEMBA1003799	0. 70
	HEMBA1003803	F-HEMBA1003803	0. 87
15	HEMBA1003804	F-HEMBA1003804	0. 94
	HEMBA1003805	F-HEMBA1003805	0. 94
	HEMBA1003807	F-HEMBA1003807,	
	HEMBA1003827	F-HEMBA1003827	0. 62
20	HEMBA1003836	F-HEMBA1003836	0. 69
	HEMBA1003838	F-HEMBA1003838	0. 80
	HEMBA1003864	F-HEMBA1003864	0, 94
	HEMBA1003866	F-HEMBA1003866	0. 32
25	HEMBA1003879	F-HEMBA1003879	0. 82
20	HEMBA1003880	F-HEMBA1003880	0. 59
	HEMBA1003893	F-HEMBA1003893	0. 47
	HEMBA1003953	F-HEMBA1003953	0. 77
00	HEMBA1003959	F-HEMBA1003959	0. 32
30	HEMBA1003978	F-HEMBA1003978	0. 37
	HEMBA1003985	F-HEMBA1003985	0.31
	HEMBA1003989	F-HEMBA1003989	0. 75
	HEMBA1004011	F-HEMBA1004011	0.34
35	HEMBA1004048	F-HEMBA1004048	0. 32
	HEMBA1004055	F-HEMBA1004055	0. 74
	HEMBA1004056	F-HEMBA1004056	0. 49
	HEMBA1004074	F-HEMBA1004074	0. 39
40	HEMBA1004086	F-HEMBA1004086	0. 45
	HEMBA1004097	F-HEMBA1004097	0. 82
	HEMBA1004111	F-HEMBA1004111	0. 94
	HEMBA1004131	F-HEMBA1004131	0. 84
45	HEMBA1004133	F-HEMBA1004133	0. 40
	HEMBA1004143	F-HEMBA1004143	0. 33
	HEMBA1004146	F-HEMBA1004146	0. 31
	HEMBA1004150	F-HEMBA1004150	0. 70
50	HEMBA1004168	F-HEMBA1004168	0. 34
	HEMBA1004199	F-HEMBA1004199	0. 34
	HEMBA1004200	F-HEMBA1004200	0.40
	HEMBA1004202	F-HEMBA1004202	0.88
55	HEMBA1004203	F-HEMBA1004203	0. 94
	HEMBA1004207	F-HEMBA1004207	0. 53

	HEMBA1004227	F-HEMBA1004227	0.94
	HEMBA1004238	F-HEMBA1004238	0. 44
5	HEMBA1004246	F-HEMBA1004246	0.71
	HEMBA1004248	F-HEMBA1004248	0. 40
	HEMBA1004264	F-HEMBA1004264	0.94
	HEMBA1004274	F-HEMBA1004274	0.88
	HEMBA1004275	F-HEMBA1004275	0. 85
10	HEMBA1004276	F-HEMBA1004276	0. 32
	HEMBA1004286	F-HEMBA1004286	0.37
	HEMBA1004289	F-HEMBA1004289	0.76
	HEMBA1004321	F-HEMBA1004321	0. 82
15	HEMBA1004327	F-HEMBA1004327	0. 40
	HEMBA1004330	F-HEMBA1004330	0.84
	HEMBA1004335	F-HEMBA1004335,	
	HEMBA1004341	F-HEMBA1004341	0. 38
20	HEMBA1004353	F-HEMBA1004353	0. 79
	HEMBA1004372	F-HEMBA1004372	0.94
	HEMBA1004389	F-HEMBA1004389	0. 32
	HEMBA1004408	F-HEMBA1004408	0.70
25	HEMBA1004479	F-HEMBA1004479	0. 55
	HEMBA1004499	F-HEMBA1004499	0.62
	HEMBA1004502	F-HEMBA1004502	0.31
	HEMBA1004507	F-HEMBA1004507	0.71
30	HEMBA1004509	F-HEMBA1004509	0.68
	HEMBA1004534	F-HEMBA1004534	0. 56
	HEMBA1004542	F-HEMBA1004542	0. 36
	HEMBA1004554	F-HEMBA1004554	0. 90
35	HEMBA1004560	F-HEMBA1004560	0. 57
	HEMBA1004573	F-HEMBA1004573	0. 93
	HEMBA1004596	F-HEMBA1004596	0. 45
	HEMBA1004604	F-HEMBA1004604	0. 91
40	HEMBA1004610	F-HEMBA1004610	0. 59
	HEMBA1004632	F-HEMBA1004632	0. 59
	HEMBA1004637	F-HEMBA1004637	0. 38
	HEMBA1004638	F-HEMBA1004638	0.40
45	HEMBA1004669	F-HEMBA1004669	0. 94
40	HEMBA1004693	F-HEMBA1004693	0. 52
	HEMBA1004697	F-HEMBA1004697	0.94
	HEMBA1004705	F-HEMBA1004705	0.94
	HEMBA1004709	F-HEMBA1004709	0.77
50	HEMBA1004711	F-HEMBA1004711	0. 73
	HEMBA1004725	F-HEMBA1004725	0. 94
	HEMBA1004734	F-HEMBA1004734	0. 94
	HEMBA1004736	F-HEMBA1004736	0. 94
55	HEMBA1004751	F-HEMBA1004751	0.64
	HEMBA1004752	F-HEMBA1004752	0. 87

	HEMBA1004753	F-HEMBA1004753 ·	0. 54
	HEMBA1004756	F-HEMBA1004756	
5	HEMBA1004758	F-HEMBA1004758	0. 69
,	HEMBA1004763	F-HEMBA1004763	0. 94
	HEMBA1004768	F-HEMBA1004768	0. 66
	HEMBA1004771	F-HEMBA1004771	0. 72
	HEMBA1004776	F-HEMBA1004776	0. 49
10	HEMBA1004795	F-HEMBA1004795	0. 46
	HEMBA1004806	F-HEMBA1004806	0. 32
	HEMBA1004847	F-HEMBA1004847	0. 79
	HEMBA1004850	F-HEMBA1004850	0. 94
15	HEMBA1004863	F-HEMBA1004863	0. 32
	HEMBA1004864	F-HEMBA1004864	0. 94
	HEMBA1004889	F-HEMBA1004889,	0. 94
	HEMBA1004923	F-HEMBA1004923	0. 70
20	HEMBA1004929	F-HEMBA1004929	0.31
	HEMBA1004930	F-HEMBA1004930	0. 90
	HEMBA1004933	F-HEMBA1004933	0.84
	HEMBA1004934	F-HEMBA1004934	0.34
25	HEMBA1004944	F-HEMBA1004944	0. 40
	HEMBA1004954	F-HEMBA1004954	0.62
	HEMBA1004972	F-HEMBA1004972	0. 90
	HEMBA1004973	F-HEMBA1004973	0.94
30	HEMBA1004977	F-HEMBA1004977	0.77
	HEMBA1004980	F-HEMBA1004980	0. 33
	HEMBA1005009	F-HEMBA1005009	0.94
	HEMBA1005019	F-HEMBA1005019	0. 32
	HEMBA1005029	F-HEMBA1005029	0.84
35	HEMBA1005035	F-HEMBA1005035	0. 91
	HEMBA1005047	F-HEMBA1005047	0. 94
	HEMBA1005050	F-HEMBA1005050	0. 88
	HEMBA1005066	F-HEMBA1005066	0. 79
40	HEMBA1005075	F-HEMBA1005075	0.31
	HEMBA1005079	F-HEMBA1005079	0. 35
	HEMBA1005083	F-HEMBA1005083	0. 43
	HEMBA1005101	F-HEMBA1005101	0. 94
45	HEMBA1005113	F-HEMBA1005113	0. 38
	HEMBA1005133	F-HEMBA1005133	0. 34
	HEMBA1005149	F-HEMBA1005149	0. 36
	HEMBA1005185	F-HEMBA1005185	0. 64
50	HEMBA1005201	F-HEMBA1005201	0. 83
	HEMBA1005202	F-HEMBA1005202	0. 94
	HEMBA1005206	F-HEMBA1005206	0. 94
	HEMBA1005219	F-HEMBA1005219	0. 57
55	HEMBA1005223	F-HEMBA1005223	0. 33
	HEMBA1005244	F-HEMBA1005244	0. 35

	HEMBA1005252	F-HEMBA1005252	0. 80
	HEMBA1005296	F-HEMBA1005296	0. 94
	HEMBA1005314	F-HEMBA1005314	0. 94
5	HEMBA1005331	F-HEMBA1005331	0. 69
	HEMBA1005338	F-HEMBA1005338	0, 94
	HEMBA1005367	F-HEMBA1005367	0. 50
	HEMBA1005382	F-HEMBA1005382	0. 91
10	HEMBA1005394	F-HEMBA1005394	0. 45
	HEMBA1005403	F-HEMBA1005403	0. 94
	HEMBA1005411	F-HEMBA1005411	0. 33
	HEMBA1005423	F-HEMBA1005423	0. 64
15	HEMBA1005468	F-HEMBA1005468	0. 33
	HEMBA1005469	F-HEMBA1005469	0. 59
	HEMBA1005472	F-HEMBA1005472,	
	HEMBA1005474	F-HEMBA1005474	0, 51
20	HEMBA1005475	F-HEMBA1005475	0. 36
20	HEMBA1005513	F-HEMBA1005513	0. 94
	HEMBA1005517	F-HEMBA1005517	0. 94
	HEMBA1005518	F-HEMBA1005518	0. 39
ar.	HEMBA1005526	F-HEMBA1005526	0. 62
25	HEMBA1005528	F-HEMBA1005528	0. 81
	HEMBA1005530	F-HEMBA1005530	0. 62
	HEMBA1005548	F-HEMBA1005548	0. 94
	HEMBA1005558	F-HEMBA1005558	0. 94
30	HEMBA1005576	F-HEMBA1005576	0. 49
	HEMBA1005581	F-HEMBA1005581	0. 53
	HEMBA1005582	F-HEMBA1005582	0.34
	HEMBA1005583	F-HEMBA1005583	0. 94
35	HEMBA1005595	F-HEMBA1005595	0. 93
	HEMBA1005609	F-HEMBA1005609	0.34
	HEMBA1005621	F-HEMBA1005621	0. 94
	HEMBA1005666	F-HEMBA1005666	0. 57
40	HEMBA1005680	F-HEMBA1005680	0. 38
	HEMBA1005685	F-HEMBA1005685	0. 32
	HEMBA1005732	F-HEMBA1005732	0. 33
	HEMBA1005737	F-HEMBA1005737	0. 92
45	HEMBA1005746	F-HEMBA1005746	0. 37
	HEMBA1005755	F-HEMBA1005755	0. 94
	HEMBA1005813	F-HEMBA1005813	0. 33
	HEMBA1005815	F-HEMBA1005815	0.82
50	HEMBA1005822	F-HEMBA1005822	0. 83
	HEMBA1005834	F-HEMBA1005834	0. 70
	HEMBA1005852	F-HEMBA1005852	0. 76
	HEMBA1005884	F-HEMBA1005884	0. 79
55	HEMBA1005891	F-HEMBA1005891	0. 94
	HEMBA1005963	F-HEMBA1005963	0. 40

	HEMBA1005990	F-HEMBA1005990	0. 82
	HEMBA1005991	F-HEMBA1005991	0. 43
	HEMBA1006031	F-HEMBA1006031	0. 55
5	HEMBA1006035	F-HEMBA1006035	0. 43
	HEMBA1006081	F-HEMBA1006081	0. 36
	HEMBA1006091	F-HEMBA1006091	0. 81
	HEMBA1006100	F-HEMBA1006100	0. 31
10	HEMBA1006108	F-HEMBA1006108	0. 80
	HEMBA1006121	F-HEMBA1006121	0. 45
	HEMBA1006130	F-HEMBA1006130	0. 46
	HEMBA1006138	F-HEMBA1006138	0. 48
15	HEMBA1006155	F-HEMBA1006155	0. 72
	HEMBA1006173	F-HEMBA1006173	0. 42
	HEMBA1006182	F-HEMBA1006182,	0. 73
	HEMBA1006198	F-HEMBA1006198	0. 94
20	HEMBA1006235	F-HEMBA1006235	0. 58
	HEMBA1006248	F-HEMBA1006248	0. 57
	HEMBA1006252	F-HEMBA1006252	0. 31
	HEMBA1006272	F-HEMBA1006272	0. 54
25	HEMBA1006278	F-HEMBA1006278	0. 92
	HEMBA1006283	F-HEMBA1006283	0.90
	HEMBA1006284	F-HEMBA1006284	0. 93
	HEMBA1006291	F-HEMBA1006291	0.38
30	HEMBA1006293	F-HEMBA1006293	0.89
30	HEMBA1006309	F-HEMBA1006309	0. 94
	HEMBA1006310	F-HEMBA1006310	0. 67
	HEMBA1006344	F-HEMBA1006344	0. 94
25	HEMBA1006347	F-HEMBA1006347	0. 94
35	HEMBA1006349	F-HEMBA1006349	0. 56
	HEMBA1006377	F-HEMBA1006377	0. 94
	HEMBA1006381	F-HEMBA1006381	0. 33
	HEMBA1006398	F-HEMBA1006398	0.71
40	HEMBA1006424		0. 40
	HEMBA1006445	F-HEMBA1006445	0. 86
	HEMBA1006467	F-HEMBA1006467	0. 55
	HEMBA1006474 HEMBA1006483	F-HEMBA1006474	0.66
45	HEMBA1006485	F-HEMBA1006483	0. 71
	HEMBA1006492	F-HEMBA1006485	0. 87 0. 73
	HEMBA1006494	F-HEMBA1006492 F-HEMBA1006494	0. 73
	HEMBA1006497	F-HEMBA1006497	0. 90
50	HEMBA1006502	F-HEMBA1006502	0. 77
	HEMBA1006507	F-HEMBA1006507	0. 56
	HEMBA1006521	F-HEMBA1006521	0. 91
	HEMBA1006530	F-HEMBA1006530	0. 37
55	HEMBA1006535	F-HEMBA1006535	0.56
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	HEMBA1006559	F-HEMBA1006559	0. 39
	HEMBA1006566	F-HEMBA1006566	0.41
	HEMBA1006579	F-HEMBA1006579	0. 83
5	HEMBA1006583	F-HEMBA1006583	
	HEMBA1006612	F-HEMBA1006612	0. 55
	HEMBA1006624	F-HEMBA1006624	0. 94
	HEMBA1006631	F-HEMBA1006631	0.74
10	HEMBA1006643	F-HEMBA1006643	0. 38
	HEMBA1006652	F-HEMBA1006652	0. 93
	HEMBA1006659	F-HEMBA1006659	0.63
	HEMBA1006674	F-HEMBA1006674	0. 54
15	HEMBA1006682	F-HEMBA1006682	0.34
	HEMBA1006708	F-HEMBA1006708	0.40
	HEMBA1006709	F-HEMBA1006709,	0. 41
	HEMBA1006717	F-HEMBA1006717	0.43
20	HEMBA1006737	F-HEMBA1006737	0. 72
	HEMBA1006754	F-HEMBA1006754	0. 45
	HEMBA1006758	F-HEMBA1006758	0.60
	HEMBA1006767	F-HEMBA1006767	0.61
25	HEMBA1006789	F-HEMBA1006789	0. 34
	HEMBA1006795	F-HEMBA1006795	0. 48
	HEMBA1006796	F-HEMBA1006796	0. 41
	HEMBA1006807	F-HEMBA1006807	0.34
30	HEMBA1006832	F-HEMBA1006832	0. 59
	HEMBA1006877	F-HEMBA1006877	0. 90
	HEMBA1006885	F-HEMBA1006885	0. 80
	HEMBA1006900	F-HEMBA1006900	0.34
35	HEMBA1006914	F-HEMBA1006914	0. 78
35	HEMBA1006926	F-HEMBA1006926	0. 67
	HEMBA1006936	F-HEMBA1006936	0. 65
	HEMBA1006941	F-HEMBA1006941	0.94
40	HEMBA1006973 HEMBA1006976	F-HEMBA1006973 F-HEMBA1006976	0. 64 0. 40
40	HEMBA1006993	F-HEMBA1006993	0. 40
	HEMBA1006996	F-HEMBA1006996	0. 77
	HEMBA1007002	F-HEMBA1007002	0. 77
45	HEMBA1007018	F-HEMBA1007018	0. 72
45	HEMBA1007052	F-HEMBA1007052	0. 41
	HEMBA1007062	F-HEMBA1007062	0. 39
	HEMBA1007066	F-HEMBA1007066	0. 35
	HEMBA1007080	F-HEMBA1007080	0. 43
50	HEMBA1007085	F-HEMBA1007085	0. 73
	HEMBA1007087	F-HEMBA1007087	0. 94
	HEMBA1007112	F-HEMBA1007112	0. 49
	HEMBA1007121	F-HEMBA1007121	0. 70
55	HEMBA1007149	F-HEMBA1007149	0. 94

	HEMBA1007151	F-HEMBA1007151	0. 80
	HEMBA1007174	F-HEMBA1007174	0. 63
	HEMBA1007178	F-HEMBA1007178	0. 47
5	HEMBA1007194	F-HEMBA1007194	0.94
	HEMBA1007203	F-HEMBA1007203	0.48
	HEMBA1007224	F-HEMBA1007224	0.74
	HEMBA1007243	F-HEMBA1007243	0. 94
10	HEMBA1007251	F-HEMBA1007251	0. 79
	HEMBA1007267	F-HEMBA1007267	0. 86
	HEMBA1007281	F-HEMBA1007281	0.61
	HEMBA1007300	F-HEMBA1007300	0. 87
15	HEMBA1007301	F-HEMBA1007301	0. 38
	HEMBA1007319	F-HEMBA1007319	0. 47
	HEMBA1007320	F-HEMBA1007320,	0. 69
	HEMBA1007342	F-HEMBA1007342	0. 80
20	HEMBB1000008	F-HEMBB1000008	0. 39
	HEMBB1000018	F-HEMBB1000018	0. 79
	HEMBB1000024	F-HEMBB1000024	0. 62
	HEMBB1000025	F-HEMBB1000025	0.66
25	HEMBB1000030	F-HEMBB1000030	0. 33
	HEMBB1000036	F-HEMBB1000036	0. 75
	HEMBB1000037	F-HEMBB1000037	0. 94
	HEMBB1000048	F-HEMBB1000048	0. 87
30	HEMBB1000083	F-HEMBB1000083	0. 48
	HEMBB1000103	F-HEMBB1000103	0. 63
	HEMBB1000119	F-HEMBB1000119	0. 94
	HEMBB1000136	F-HEMBB1000136	0. 92
35	HEMBB1000173	F-HEMBB1000173	0. 46
	HEMBB1000175	F-HEMBB1000175	0. 48
	HEMBB1000198	F-HEMBB1000198	0. 74
	HEMBB1000215	F-HEMBB1000215	0. 61
40	HEMBB1000217	F-HEMBB1000217	0. 94
40	HEMBB1000226 HEMBB1000240	F-HEMBB1000226	0. 94
	HEMBB1000240	F-HEMBB1000240	0. 40
	HEMBB1000244	F-HEMBB1000244	0.81
	HEMBB1000200	F-HEMBB1000266 F-HEMBB1000338	0. 92
45	HEMBB1000338	F-HEMBB1000339	0. 33 0. 40
	HEMBB1000339	F-HEMBB1000391	0. 40
	HEMBB1000331	F-HEMBB1000438	0. 79
	HEMBB1000438	F-HEMBB1000438	0. 79
50	HEMBB1000510	F-HEMBB1000510	0. 35
	HEMBB1000550	F-HEMBB1000550	0. 94
	HEMBB1000556	F-HEMBB1000556	0. 89
	HEMBB1000589	F-HEMBB1000589	0. 77
55	HEMBB1000591	F-HEMBB1000591	0. 77
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	HEMBB1000592	F-HEMBB1000592	0.77
	HEMBB1000593	F-HEMBB1000593	0.68
	HEMBB1000623	F-HEMBB1000623	0.94
5	HEMBB1000630	F-HEMBB1000630	0.59
	HEMBB1000631	F-HEMBB1000631	0.71
	HEMBB1000632	F-HEMBB1000632	0.38
	HEMBB1000671	F-HEMBB1000671	0.36
10	HEMBB1000673	F-HEMBB1000673	0. 54
	HEMBB1000693	F-HEMBB1000693	0.68
	HEMBB1000705	F-HEMBB1000705	0.44
	HEMBB1000706	F-HEMBB1000706	0.45
15	HEMBB1000725	F-HEMBB1000725	0.91
	HEMBB1000763	F-HEMBB1000763	0.87
	HEMBB1000774	F-HEMBB1000774,	0.80
	HEMBB1000781	F-HEMBB1000781	0.94
20	HEMBB1000789	F-HEMBB1000789	0.34
	HEMBB1000807	F-HEMBB1000807	0.94
	HEMBB1000810	F-HEMBB1000810	0.33
	HEMBB1000826	F-HEMBB1000826	0.36
25	HEMBB1000835	F-HEMBB1000835	0.81
	HEMBB1000848	F-HEMBB1000848	0.66
	HEMBB1000852	F-HEMBB1000852	0.85
	HEMBB1000870	F-HEMBB1000870	0.34
30	HEMBB1000887	F-HEMBB1000887	0. 58 0. 31
	HEMBB1000908 HEMBB1000927	F-HEMBB1000908 F-HEMBB1000927	0. 94
	HEMBB1000947	F-HEMBB1000947	0.33
	HEMBB1000973	F-HEMBB1000973	0. 72
35	HEMBB1000975	F-HEMBB1000975	0.68
	HEMBB1000985	F-HEMBB1000985	0. 62
	HEMBB1000991	F-HEMBB1000991	0. 45
	HEMBB1001011	F-HEMBB1001011	0.58
40	HEMBB1001014	F-HEMB81001014	0.86
	HEMBB1001020	F-HEMBB1001020	0. 87
	HEMBB1001024	F-HEMBB1001024	0.64
	HEMBB1001056	F-HEMBB1001056	0. 94
45	HEMBB1001058	F-HEMBB1001058	0. 39
	HEMBB1001068	F-HEMBB1001068	0.57
	HEMBB1001096	F-HEMBB1001096	0. 48
	HEMBB1001105	F-HEMBB1001105	0. 45
50	HEMBB1001112	F-HEMBB1001112	0.47
	HEMBB1001117	F-HEMBB1001117	0.83
	HEMBB1001126	F-HEMBB1001126	0.76
	HEMBB1001137	F-HEMBB1001137	0.53
55	HEMBB1001151	F-HEMBB1001151	0.94
	HEMBB1001153	F-HEMBB1001153	0. 77

	HEMBB1001169	F-HEMBB1001169	0. 39
	HEMBB1001175	F-HEMBB1001175	0. 91
	HEMBB1001182	F-HEMBB1001182	0. 94
5	HEMBB1001199	F-HEMBB1001199	0. 54
	HEMBB1001210	F-HEMBB1001210	0. 88
	HEMBB1001234	F-HEMBB1001234	0. 67
	HEMBB1001242	F-HEMBB1001242	0. 33
10	HEMB81001288	F-HEMBB1001288	0. 33
	HEMBB1001289	F-HEMBB1001289	0. 58
	HEMBB1001294	F-HEMB81001294	0. 86
	HEMBB1001314	F-HEMBB1001314	0. 76
15	HEMBB1001331	F-HEMBB1001331	0. 44
	HEMBB1001339	F-HEMBB1001339	0.40
	HEMBB1001346	F-HEMBB1001346,	0. 83
	HEMBB1001364	F-HEMBB1001364	0. 69
20	HEMBB1001369	F-HEMBB1001369	0. 47
	HEMBB1001384	F-HEMBB1001384	0.80
	HEMBB1001387	F-HEMBB1001387	0. 36
	HEMBB1001394	F-HEMBB1001394	0. 94
25	HEMBB1001410	F-HEMBB1001410	0.51
	HEMBB1001426	F-HEMBB1001426	0. 80
	HEMBB1001429	F-HEMBB1001429	0. 91
	HEMBB1001449	F-HEMBB1001449	0. 31
30	HEMBB1001482	F-HEMBB1001482	0. 44
	HEMBB1001531	F-HEMBB1001531	0. 58
	HEMBB1001562	F-HEMBB1001562	0. 65
	HEMBB1001564	F-HEMB81001564	0. 46
35	HEMBB1001585	F-HEMBB1001585	0. 74
-	HEMBB1001603	F-HEMBB1001603	0. 81
	HEMBB1001635	F-HEMBB1001635	0. 46
	HEMBB1001653	F-HEMBB1001653	0. 33
40	HEMBB1001665	F-HEMBB1001665	0. 39
40	HEMBB1001668	F-HEMBB1001668 F-HEMBB1001673	0. 38
	HEMBB1001673 HEMBB1001684		0. 83 0. 44
	HEMBB1001685	F-HEMBB1001684 F-HEMBB1001685	
	HEMBB1001695	F-HEMBB1001695	0. 90 0. 92
45	HEMBB1001707	F-HEMBB1001707	0. 52
	HEMBB1001736	F-HEMBB1001736	0. 94
	HEMBB1001760	F-HEMBB1001760	0. 94
	HEMBB1001785	F-HEMBB1001785	0. 74
50	HEMBB1001812	F-HEMBB1001812	0. 74
	HEMBB1001816	F-HEMBB1001816	0. 33
	HEMBB1001831	F-HEMBB1001831	0. 41
	HEMBB1001834	F-HEMBB1001834	0. 93
55	HEMBB1001839	F-HEMBB1001839	0. 78
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	HEMBB1001869	F-HEMBB1001869	0. 48
	HEMBB1001872	F-HEMBB1001872	0. 93
	HEMBB1001874	F-HEMBB1001874	0. 31
5	HEMBB1001905	F-HEMBB1001905	0. 87
	HEMBB1001906	F-HEMBB1001906	0. 88
	HEMBB1001908	F-HEMBB1001908	0. 69
	HEMBB1001910	F-HEMBB1001910	0. 77
10	HEMBB1001915	F-HEMBB1001915	0. 76
	HEMBB1001945	F-HEMBB1001945	0. 64
	HEMBB1001947	F-HEMBB1001947	0. 94
	HEMBB1001950	F-HEMBB1001950	0. 73
15	HEMBB1001957	F-HEMBB1001957	0. 64
	HEMBB1001962	F-HEMBB1001962	0. 90
	HEMBB1001990	F-HEMBB1001990,	0. 83
	HEMBB1002044	F-HEMBB1002044	0. 82
20	HEMBB1002050	F-HEMBB1002050	0. 91
20	HEMBB1002068	F-HEMBB1002068	0. 48
	HEMBB1002134	F-HEMBB1002134	0. 80
	HEMBB1002142	F-HEMBB1002142	0. 39
	HEMBB1002152	F-HEMBB1002152	0. 31
25	HEMBB1002193	F-HEMBB1002193	0. 35
	HEMBB1002217	F-HEMBB1002217	0. 49
	HEMBB1002218	F-HEMBB1002218	0. 32
	HEMBB1002249	F-HEMBB1002249	0. 72
30	HEMBB1002300	F-HEMBB1002300	0. 75
	HEMBB1002327	F-HEMBB1002327	0.86
	HEMBB1002329	F-HEMBB1002329	0. 31
	HEMBB1002340	F-HEMBB1002340	0. 42
35	HEMBB1002342	F-HEMBB1002342	0. 76
	HEMBB1002358	F-HEMBB1002358	0.70
	HEMBB1002359	F~HEMBB1002359	0. 75
	HEMBB1002383	F-HEMBB1002383	0. 78
40	HEMBB1002409	F-HEMBB1002409	0. 52
	HEMBB1002415	F-HEMBB1002415	0.82
	HEMBB1002442	F-HEMBB1002442	0. 31
	HEMBB1002457	F-HEMBB1002457	0. 46
45	HEMBB1002489	F-HEMBB1002489	0. 69
	HEMBB1002492	F-HEMBB1002492	0. 68
	HEMBB1002495	F-HEMBB1002495	0.94
	HEMBB1002502	F-HEMBB1002502	0. 39
50	HEMBB1002510	F-HEMB81002510	0. 94
	HEMBB1002545	F-HEMBB1002545	0. 44
	HEMBB1002550	F-HEMBB1002550	0. 57
	HEMBB1002600	F-HEMBB1002600	0. 91
55	HEMBB1002607	F-HEMBB1002607	0. 83
55	HEMBB1002614	F-HEMBB1002614	0. 75

	HEMBB1002684	F-HEMBB1002684	0. 84
	HEMBB1002692	F-HEMBB1002692	0. 52
	HEMBB1002697	F-HEMBB1002697	0. 86
5	HEMBB1002705	F-HEMBB1002705	0. 32
	MAMMA1000019	F-MAMMA1000019	0. 48
	MAMMA1000020	F-MAMMA1000020	0.36
	MAMMA1000025	F-MAMMA1000025	0. 31
10	MAMMA1000055	F-MAMMA1000055	0. 39
	MAMMA1000057	F-MAMMA1000057	0. 35
	MAMMA1000069	F-MAMMA1000069	0.78
	MAMMA1000084	F-MAMMA1000084	0.61
15	MAMMA1000085	F-MAMMA1000085	0.72
	MAMMA1000139	F-MAMMA1000139	0. 33
	MAMMA1000143	F-MAMMA1000143,	0.40
	MAMMA1000163	F-MAMMA1000163	0. 94
20	MAMMA1000171	F-MAMMA1000171	0. 78
	MAMMA1000173	F-MAMMA1000173	0. 94
	MAMMA1000183	F-MAMMA1000183	0.77
	MAMMA1000251	F-MAMMA1000251	0.94
25	MAMMA1000257	F-MAMMA1000257	0. 75
	MAMMA1000277	F-MAMMA1000277	0.94
	MAMMA1000278	F-MAMMA1000278	0. 35
	MAMMA1000279	F-MAMMA1000279	0.62
30	MAMMA1000284	F-MAMMA1000284	0. 35
30	MAMMA1000309	F-MAMMA1000309	0. 53
	MAMMA1000312	F-MAMMA1000312	0. 68
	MAMMA1000313	F-MAMMA1000313	0. 32
	MAMMA1000339	F-MAMMA1000339	0. 35
35	MAMMA1000361	F-MAMMA1000361	0. 61
	MAMMA1000372	F-MAMMA1000372	0. 60
	MAMMA1000388	F-MAMMA1000388	0. 93
	MAMMA1000395	F-MAMMA1000395	0. 94
40	MAMMA1000410	F-MAMMA1000410	0. 94
	MAMMA1000416	F-MAMMA1000416	0. 89
	MAMMA1000421	F-MAMMA1000421	0.50
	MAMMA1000422		0. 92
45	MAMMA1000429	F-MAMMA1000429	0. 92
	MAMMA1000458	F-MAMMA1000458	0. 88
	MANMA1000468	F-MAMMA1000468	0.60
	MAMMA1000472	F-MAMMA1000472	0. 33
50	MAMMA1000490 MAMMA1000524	F-MAMMA1000490	0.34
	MAMMA1000567	F-MAMMA1000524	0.39
	MAMMA1000583	F-MAMMA1000567 F-MAMMA1000583	0.40
	MAMMA1000612	F-MAMMA1000612	0.45
55	MAMMA1000612	F-MAMMA1000612	0.89
	medianes i UUUU43	r mamma 1000023	0. 44

	MAMMA1000625	F-MAMMA1000625	0. 52
	MAMMA1000664	F-MAMMA1000664	0. 42
	MAMMA1000670	F-MAMMA1000670	0. 94
5	MAMMA1000672	F-MAMMA1000672	0.44
	MAMMA1000684	F-MAMMA1000684	0.94
	MAMMA1000713	F-MAMMA1000713	0.88
	MAMMA1000731	F-MAMMA1000731	0. 67
10	MAMMA1000734	F-MAMMA1000734	0.40
	MAMMA1000738	F-MAMMA1000738	0. 93
	MAMMA1000746	F-MAMMA1000746	0.48
	MAMMA1000760	F-MAMMA1000760	0.94
15	MAMMA1000776	F-MAMMA1000776	0. 94
	MAMMA1000824	F-MAMMA1000824	0. 81
	MAMMA1000831	F-MAMMA1000831	0.39
	MAMMA1000841	F-MAMMA1000841	0.41
20	MAMMA1000842	F-MAMMA1000842	0. 61
	MAMMA1000843	F-MAMMA1000843	0. 48
	MAMMA1000856	F-MAMMA1000856	0.31
	MAMMA1000859	F-MAMMA1000859	0. 44
25	MAMMA1000865	F-MAMMA1000865	0. 94
	MAMMA1000875	F-MAMMA1000875	0. 49
	MAMMA1000897	F-MAMMA1000897	0.94
	MAMMA1000906	F-MAMMA1000906	0. 45
30	MAMMA1000908	F-MAMMA1000908	0. 39
	MAMMA1000914	F-MAMMA1000914	0.94
	MAMMA1000921	F-MAMMA1000921	0. 34
	MAMMA1000956	F-MAMMA1000956	0.39
35	MAMMA1000968 MAMMA1000979	F-MAMMA1000968	0. 35
	MAMMA1000979	F-MAMMA1000979 F-MAMMA1001008	0. 68 0. 36
	MAMMA1001003	F-MAMMA1001008	0. 48
	MAMMA1001041	F-MAMMA1001041	0. 34
40	MAMMA1001059	F-MAMMA1001059	0. 94
	MAMMA1001073	F-MAMMA1001073	0. 94
	MAMMA1001075	F-MAMMA1001075	0. 94
	MAMMA1001078	F-MAMMA1001078	0. 52
45	MAMMA1001080	F-MAMMA1001080	0. 87
	MAMMA1001091	F-MAMMA1001091	0. 53
	MAMMA1001105	F-MAMMA1001105	0. 94
	MAMMA1001110	F-MAMMA1001110	0. 78
50	MAMMA1001126	F-MAMMA1001126	0. 85
	MAMMA1001139	F-MAMMA1001139	0. 38
	MAMMA1001143	F-MAMMA1001143	0. 79
	MAMMA1001154	F-MAMMA1001154	0. 31
55	MAMMA1001181	F-MAMMA1001181	0. 69
	MAMMA1001198	F-MAMMA1001198	0. 88

	MAMMA1001202	F-MAMMA1001202	0. 38
	MAMMA1001215	F-MAMMA1001215	0. 94
5	MAMMA1001222	F-MAMMA1001222	0.89
Ť	MAMMA1001244	F-MAMMA1001244	0. 87
	MAMMA1001259	F-MAMMA1001259	0. 91
	MAMMA1001260	F-MAMMA1001260	0.75
	MAMMA1001271	F-MAMMA1001271	0.77
10	MAMMA1001292	F-MAMMA1001292	0.94
	MAMMA1001296	F-MAMMA1001296	0.75
	MAMMA1001305	F-MAMMA1001305	0.66
	MAMMA1001343	F-MAMMA1001343	0.38
15	MAMMA1001346	F-MAMMA1001346	0.81
	MAMMA1001388	F-MAMMA1001388	0.94
	MAMMA1001411	F-MAMMA1001411,	0. 42
	MAMMA1001419	F-MAMMA1001419	0. 45
20	MAMMA1001465	F-MAMMA1001465	0.94
	MAMMA1001476	F-MAMMA1001476	0. 90
	MAMMA1001487	F-MAMMA1001487	0. 78
	MAMMA1001510	F-MAMMA1001510	0.82
25	MAMMA1001522	F-MAMMA1001522	0.63
	MAMMA1001551	F-MAMMA1001551	0.81
	MAMMA1001576	F-MAMMA1001576	0.94
	MAMMA1001600	F-MAMMA1001600	0. 72
30	MAMMA1001604	F-MAMMA1001604	0. 87
30	MAMMA1001620	F-MAMMA1001620	0.43
	MAMMA1001627	F-MAMMA1001627	0.39
	MAMMA1001630	F-MAMMA1001630	0.46
	MAMMA1001633	F-MAMMA1001633	0.58
35	MAMMA1001635	F-MAMMA1001635	0.89
	MAMMA1001649	F-MAMMA1001649	0.62
	MAMMA1001654	F-MAMMA1001654	0.94
	MAMMA1001692	F-MAMMA1001692	0.83
40	MAMMA1001730	F-MAMMA1001730	0.48
	MAMMA1001735	F-MAMMA1001735	0.94
	MAMMA1001743	F-MAMMA1001743	0.94
	MAMMA1001744	F-MAMMA1001744	0. 33
45	MAMMA1001751	F-MAMMA1001751	0.94
	MAMMA1001754	F-MAMMA1001754	0. 47
	MAMMA1001757	F-MAMMA1001757	0.38
	MAMMA1001764	F-MAMMA1001764	0.57
50	MAMMA1001768	F-MAMMA1001768	0. 85
	MAMMA1001771	F-MAMMA1001771	0.39
	MAMMA1001783	F-MAMMA1001783	0.61
	MAMMA1001785	F-MAMMA1001785	0.41
55	MAMMA1001790	F-MAMMA1001790	0.69
	MAMMA1001812	F-MAMMA1001812	0.32

	MAMMA1001817	F-MAMMA1001817	0. 55
	MAMMA1001824	F-MAMMA1001824	0.57
	MAMMA1001837	F-MAMMA1001837	0. 35
5	MAMMA1001848	F-MAMMA1001848	0.38
	MAMMA1001851	F-MAMMA1001851	0. 46
	MAMMA1001858	F-MAMMA1001858	0. 53
	MAMMA1001864	F-MAMMA1001864	0.32
10	MAMMA1001868	F-MAMMA1001868	0.48
	MAMMA1001874	F-MAMMA1001874	0. 94
	MAMMA1001956	F-MAMMA1001956	0. 47
	MAMMA1001969	F-MAMMA1001969	0. 31
15	MAMMA1001970	F-MAMMA1001970	0.34
	MAMMA1002009	F-MAMMA1002009	0. 45
	MAMMA1002011	F-MAMMA1002011,	0.94
	MAMMA1002033	F-MAMMA1002033	0. 33
20	MAMMA1002041	F-MAMMA1002041	0.86
	MAMMA1002042	F-MAMMA1002042	0. 88
	MAMMA1002047	F-MAMMA1002047	0. 41
	MAMMA1002068	F-MAMMA1002068	0. 38
25	MAMMA1002143	F-MAMMA1002143	0. 91
	MAMMA1002153	F-MAMMA1002153	0. 94
	MAMMA1002156	F-MAMMA1002156	0. 73
	MAMMA1002170	F-MAMMA1002170	0. 43
30	MAMMA1002174	F-MAMMA1002174	0. 69
	MAMMA1002198	F-MAMMA1002198	0. 65
	MAMMA1002209	F-MAMMA1002209	0.80
	MAMMA1002219	F-MAMMA1002219	0.85
35	MAMMA1002236 MAMMA1002243	F-MAMMA1002236 F-MAMMA1002243	0. 55 0. 50
	MAMMA1002243	F-MAMMA1002243	0. 50
	MAMMA1002269	F-MAMMA1002269	0. 43
	MAMMA1002292	F-MAMMA1002292	0. 76
40	MAMMA1002294	F-MAMMA1002294	0. 76
	MAMMA1002297	F-MAMMA1002297	0. 91
	MAMMA1002308.	F-MAMMA1002308	0. 68
	MAMMA1002312	F-MAMMA1002312	0.77
45	MAMMA1002317	F-MAMMA1002317	0.46
	MAMMA1002319	F-MAMMA1002319	0. 31
	MAMMA1002329	F-MAMMA1002329	0.34
	MAMMA1002333	F-MAMMA1002333	0. 65
50	MAMMA1002351	F-MAMMA1002351	0. 89
	MAMMA1002353	F-MAMMA1002353	0. 94
	MAMMA1002355	F-MAMMA1002355	0.55
	MAMMA1002356	F-MAMMA1002356	0.32
55	MAMMA1002362	F-MAMMA1002362	0.83
	MAMMA1002380	F-MAMMA1002380	0. 94

	MAMMA1002384 F-	WAMMA1002384	0. 42
	MAMMA1002385 F-	MAMMA1002385	0. 82
	MAMMA1002413 F-	MAMMA1002413	0. 94
5	MAMMA1002427 F-1	MAMMA1002427	
	MAMMA1002434 F-1	MAMMA1002434	0. 74
	MAMMA1002454 F-	MAMMA1002454	0. 82
		MAMMA1002461	0.49
10		MAMMA1002470	0.89
		MAMMA1002485	0. 55
	MAMMA1002524 F-	MAMMA1002524	0.33
	MAMMA1002530 F-	MAMMA1002530	0.75
15		MAMMA1002554	0. 41
		MAMMA1002585	0. 31
		MAMMA1002590,	
		MAMMA1002598	0. 76
20	MAMMA1002617 F-M	MAMMA1002617	0.49
	MAMMA1002619 F-1	MAMMA1002619	0.88
	MAMMA1002637 F-1	AAMMA1002637	0. 94
	MAMMA1002650 F-	MAMMA1002650	0. 54
25	MAMMA1002655 F-N	MAMMA1002655	0. 39
	MAMMA1002665 F-N	MAMMA1002665	0. 33
	MAMMA1002671 F-M	MAMMA1002671	0. 94
	MAMMA1002673 F-M	MAMMA1002673	0.70
30	. MAMMA1002684 F-1	MAMMA1002684	0.94
••	MAMMA1002685 F-8	MAMMA1002685	0.88
	MAMMA1002699 F-H	MAMMA1002699	0.94
	MAMMA1002711 F-N	MAMMA1002711	0.39
35	MAMMA1002769 F-N	MAMMA1002769	0.88
33	MAMMA1002775 F-H	MAMMA1002775	0.81
	MAMMA1002782 F-	MAMMA1002782	0. 87
		MAMMA1002796	0. 40
		MAMMA1002807	0. 64
40		MAMMA1002842	0. 94
	· · · · · · · · · · · · · · · · · · ·	MAMMA1002843	0. 32
		MAMMA1002868	
		MAMMA1002869	0. 94
45		MAMMA1002881	0. 94
		MAMMA1002886	0. 61
		MAMMA1002890	0. 94
		MAMMA1002895	0. 94
50		MAMMA1002937	0. 91
		MAMMA1002938	0. 78
		MAMMA1002947	0. 94
		MAMMA1002964	0. 34
55		MAMMA1002987	0. 74
	MAMMA1003011 F-N	MAMMA1003011	0. 90

	MAMMA1003013	F-MAMMA1003013	0. 36
	MAMMA1003015	F-MAMMA1003015	0. 92
E	MAMMA1003035	F-MAMMA1003035	0. 94
5	MAMMA1003044	F-MAMMA1003044	0. 61
	MAMMA1003047	F-MAMMA1003047	0. 94
	MAMMA1003049	F-MAMMA1003049	0.31
	MAMMA1003056	F-MAMMA1003056	0. 75
10	MAMMA1003057	F-MAMMA1003057	0. 33
	MAMMA1003066	F-MAMMA1003066	0. 38
	MAMMA1003099	F-MAMMA1003099	0. 79
	MAMMA1003104	F-MAMMA1003104	0. 44
15	MAMMA1003113	F-MAMMA1003113	0. 90
	MAMMA1003127	F-MAMMA1003127	0. 53
	MAMMA1003135	F-MAMMA1003135,	0. 94
	MAMMA1003146	F-MAMMA1003146	0. 88
20	MAMMA1003150	F-MAMMA1003150	0. 73
	MAMMA1003166	F-MAMMA1003166	0. 94
	NT2RM1000001	F-NT2RM1000001	0. 66
	NT2RM1000018	F-NT2RM1000018	0. 55
25	NT2RM1000032	F-NT2RM1000032	0. 74
	NT2RM1000035	F-NT2RM1000035	0. 52
	NT2RM1000037	F-NT2RM1000037	0. 94
	NT2RM1000039	F-NT2RM1000039	0.77
30	NT2RM1000055	F-NT2RM1000055	0.89
•	NT2RM1000059	F-NT2RM1000059	0.41
	NT2RM1000062	F-NT2RM1000062	0. 65
	NT2RM1000080	F-NT2RM1000080	0.61
35	NT2RM1000086	F-NT2RM1000086	0. 94
	NT2RM1000092	F-NT2RM1000092	0. 94
	NT2RM1000118	F-NT2RM1000118	0. 76
	NT2RM1000119	F-NT2RM1000119	0. 41
40	NT2RM1000127	F-NT2RM1000127	0. 94
40	NT2RM1000131	F-NT2RM1000131	0. 90
	NT2RM1000132	F-NT2RM1000132	0. 94
	NT2RM1000153	F-NT2RM1000153	0. 90
	NT2RM1000186	F-NT2RM1000186	0. 78
45	NT2RM1000187	F-NT2RM1000187	0. 54
	NT2RM1000199	F-NT2RM1000199	0. 33
	NT2RM1000242	F-NT2RM1000242	0. 78
	NT2RM1000244	F-NT2RM1000244	0.31
50	NT2RM1000252	F-NT2RM1000252	0.76
	NT2RM1000256	F-NT2RM1000256	0. 94
	NT2RM1000257	F-NT2RM1000257	0.89
	NT2RM1000260	F-NT2RM1000260	0. 42
55	NT2RM1000271	F-NT2RM1000271	0. 74
	NT2RM1000272	F-NT2RM1000272	0. 78

	NTODUTOOOOO	E-NTORNI 00000	0.04
	NT2RM1000280	F-NT2RM1000280	0. 94
	NT2RM1000300	F-NT2RM1000300	0. 47
5	NT2RM1000314	F-NT2RM1000314	0.61
3	NT2RM1000318	F-NT2RM1000318	• . •
	NT2RM1000341	F-NT2RM1000341	0. 70
	NT2RM1000354	F-NT2RM1000354	0. 82
	NT2RM1000355	F-NT2RM1000355	0. 94
10	NT2RM1000365	F-NT2RM1000365	0.31
	NT2RM1000377	F-NT2RM1000377	0. 87
	NT2RM1000388	F-NT2RM1000388	0.94
	NT2RM1000394	F-NT2RM1000394	0.58
15	NT2RM1000399	F-NT2RM1000399	0.86
	NT2RM1000421	F-NT2RM1000421	0.39
	NT2RM1000430	F-NT2RM1000430,	0.94
	NT2RM1000499	F-NT2RM1000499	0.44
20	NT2RM1000539	F-NT2RM1000539	0.87
	NT2RM1000553	F-NT2RM1000553	0.88
	NT2RM1000555	F-NT2RM1000555	0.37
	NT2RM1000563	F-NT2RM1000563	0.63
25	NT2RM1000623	F-NT2RM1000623	0.34
20	NT2RM1000648	F-NT2RM1000648	0.94
	NT2RM1000661	F-NT2RM1000661	0.61
	NT2RM1000666	F-NT2RM1000666	0.94
	NT2RM1000669	F-NT2RM1000669	0.94
30	NT2RM1000672	F-NT2RM1000672	0.53
	NT2RM1000691	F-NT2RM1000691	0.38
	NT2RM1000699	F-NT2RM1000699	0.32
	NT2RM1000702	F-NT2RM1000702	0.74
35	NT2RM1000725	F-NT2RM1000725	0.75
	NT2RM1000741	F-NT2RM1000741	0.94
	NT2RM1000742	F-NT2RM1000742	0.54
	NT2RM1000746	F-NT2RM1000746	0.94
40	NT2RM1000770	F-NT2RM1000770	0.48
	NT2RM1000772	F-NT2RM1000772	0.39
	NT2RM1000780	F-NT2RM1000780	0.66
	NT2RM1000781	F-NT2RM1000781	0.49
45	NT2RM1000800	F-NT2RM1000800	0.72
	NT2RM1000802	F-NT2RM1000802	0.89
	NT2RM1000811	F-NT2RM1000811	0. 53
	NT2RM1000826	F-NT2RM1000826	0.39
50	NT2RM1000829	F-NT2RM1000829	0. 63
	NT2RM1000833	F-NT2RM1000833	0.94
	NT2RM1000850	F-NT2RM1000850	0.89
	NT2RM1000852	F-NT2RM1000852	0.88
	NT2RM1000857	F-NT2RM1000857	0.58
55	NT2RM1000867	F-NT2RM1000867	0. 90

	NT2RM1000874	F-NT2RM1000874	0. 89
	NT2RM1000882	F-NT2RM1000882	0. 63
	NT2RM1000883	F-NT2RM1000883	0. 71
5	NT2RM1000885	F-NT2RM1000885	
	NT2RM1000894	F-NT2RM1000894	0. 57
	NT2RM1000898	F-NT2RM1000898	0. 87
	NT2RM1000905	F-NT2RM1000905	0. 94
10	NT2RM1000924	F-NT2RM1000924	0. 88
	NT2RM1000927	F-NT2RM1000927	0. 70
	NT2RM1000962	F-NT2RM1000962	0. 84
	NT2RM1000978	F-NT2RM1000978	0. 39
15	NT2RM1001003	F-NT2RM1001003	0. 81
	NT2RM1001008	F-NT2RM1001008	0. 94
	NT2RM1001043	F-NT2RM1001043,	0. 94
	NT2RM1001044	F-NT2RM1001044	0. 94
20	NT2RM1001059	F-NT2RM1001059	0.94
	NT2RM1001066	F-NT2RM1001066	0. 94
	NT2RM1001072	F-NT2RM1001072	0. 77
	NT2RM1001074	F-NT2RM1001074	0. 94
25	NT2RM1001082	F-NT2RM1001082	0. 94
	NT2RM1001085	F-NT2RM1001085	0.83
	NT2RM1001092	F-NT2RM1001092	0. 77
	NT2RM1001102	F-NT2RM1001102	0. 59
30	NT2RM1001105	F-NT2RM1001105	0.94
	NT2RM1001112	F-NT2RM1001112	0. 50
	NT2RM1001115	F-NT2RM1001115	0. 32
	NT2RM1001139	F-NT2RM1001139	0. 77
35	NT2RM2000006	F-NT2RM2000006	0. 34
	NT2RM2000013	F-NT2RM2000013	0. 70
	NT2RM2000030	F-NT2RM2000030	0. 89
	NT2RM2000032	F-NT2RM2000032	0. 36
40	NT2RM2000042 NT2RM2000092	F-NT2RM2000042 F-NT2RM2000092	0. 32
	NT2RM2000092	F-NT2RM2000092	0. 55 0. 33
	NT2RM2000101	F-NT2RM2000101	
	NT2RM2000124		0. 77
45	NT2RM2000191	F-NT2RM2000124	0. 80
	NT2RM2000192	F-NT2RM2000192	0. 93
	NT2RM2000239	F-NT2RM2000239	0. 68
	NNNNNNNNNNN	F-NNNNNNNNNNNN	0. 94
50	NT2RM2000250	F-NT2RM2000250	0. 94
•	NT2RM2000259	F-NT2RM2000259	0. 68
	NT2RM2000260	F-NT2RM2000260	0. 37
	NT2RM2000287	F-NT2RM2000287	0. 46
55	NT2RM2000322	F-NT2RM2000322	0. 83
55	NT2RM2000359	F-NT2RM2000359	0. 94

	NT2RM2000363	F-NT2RM2000363	0.91
	NT2RM2000368	F-NT2RM2000368	0. 34
5	NT2RM2000371	F-NT2RM2000371	0. 94
•	NT2RM20003 <u>7</u> 4	F-NT2RM2000374_	0, 65
	NT2RM2000395	F-NT2RM2000395	0. 33
	NT2RM2000402	F-NT2RM2000402	0.80
10	NT2RM2000407	F-NT2RM2000407	0.88
10	NT2RM2000420	F-NT2RM2000420	0. 92
	NT2RM2000422	F-NT2RM2000422	0.49
	NT2RM2000452	F-NT2RM2000452	0. 93
	NT2RM2000469	F-NT2RM2000469	0.39
15	NT2RM2000490	F-NT2RM2000490	0.90
	NT2RM2000502	F-NT2RM2000502	0.64
	NT2RM2000504	F-NT2RM2000504;	0.71
	NT2RM2000522	F-NT2RM2000522	0.91
20	NT2RM2000540	F-NT2RM2000540	0.87
	NT2RM2000556	F-NT2RM2000556	0.91
	NT2RM2000566	F-NT2RM2000566	0.39
	NT2RM2000567	F-NT2RM2000567	0.44
25	NT2RM2000569	F-NT2RM2000569	0. 49
	NT2RM2000577	F-NT2RM2000577	0. 45
	NT2RM2000581	F-NT2RM2000581	0. 72
	NT2RM2000588	F-NT2RM2000588	0.31
30	NT2RM2000594	F-NT2RM2000594	0. 79
	NT2RM2000599	F-NT2RM2000599	0. 80
	NT2RM2000612	F-NT2RM2000612	0. 75
	NT2RM2000623	F-NT2RM2000623	0. 72
35	NT2RM2000624	F-NT2RM2000624	0. 94
	NT2RM2000635	F-NT2RM2000635	0. 73
	NT2RM2000636	F-NT2RM2000636	0. 45
	NT2RM2000639	F-NT2RM2000639	0. 43
40	NT2RM2000649	F-NT2RM2000649	0. 74
	NT2RM2000669	F-NT2RM2000669	0. 94
	NT2RM2000691	F-NT2RM2000691	0. 84
	NT2RM2000714	F-NT2RM2000714	0.92
45	NT2RM2000718 NT2RM2000735	F-NT2RM2000718	0. 52
70		F-NT2RM2000735	0. 33
	NT2RM2000740 NT2RM2000795	F-NT2RM2000740	0. 94
		F-NT2RM2000795	0. 44
50	NT2RM2000821	F-NT2RM2000821	0. 33
50	NT2RM2000837 NT2RM2000951	F-NT2RM2000837 F-NT2RM2000951	0.57
			0.32
	NT2RM2000952	F-NT2RM2000952	0.50
	NT2RM2000984	F-NT2RM2000984	0.76
55	NT2RM2001004	F-NT2RM2001004	0.55
	NT2RM2001035	F-NT2RM2001035	0. 82

	NT2RM2001065	F-NT2RM2001065	0. 65
	NT2RM2001100	F-NT2RM2001100	0. 89
5	NT2RM2001105	F-NT2RM2001105	0. 94
·	NT2RM2001131	F-NT2RM2001131	0.94
	NT2RM2001141	F-NT2RM2001141	0.74
	NT2RM2001152	F-NT2RM2001152	0. 33
-	NT2RM2001177	F-NT2RM2001177	0. 32
10	NT2RM2001194	F-NT2RM2001194	0. 83
	NT2RM2001196	F-NT2RM2001196	0. 42
	NT2RM2001201	F-NT2RM2001201	0. 90
	NT2RM2001221	F-NT2RM2001221	0.39
15	NT2RM2001238	F-NT2RM2001238	0. 94
	NT2RM2001243	F-NT2RM2001243	0. 44
	NT2RM2001247	F-NT2RM2001247,	
	NT2RM2001256	F-NT2RM2001256	0, 60
20	NT2RM2001291	F-NT2RM2001291	0. 39
	NT2RM2001306	F-NT2RM2001306	0. 81
	NT2RM2001312	F-NT2RM2001312	0. 64
	NT2RM2001319	F-NT2RM2001319	0. 53
25	NT2RM2001324	F-NT2RM2001324	0. 94
	NT2RM2001345	F-NT2RM2001345	0. 94
	NT2RM2001360	F-NT2RM2001360	0.69
	NT2RM2001370	F-NT2RM2001370	0.44
30	NT2RM2001393	F-NT2RM2001393	0.94
	NT2RM2001420	F-NT2RM2001420	0.46
	NT2RM2001424	F-NT2RM2001424	0. 94
	NT2RM2001499	F-NT2RM2001499	0. 63
35	NT2RM2001504	F-NT2RM2001504	0. 94
33	NT2RM2001524	F-NT2RM2001524	0. 76
	NT2RM2001544	F-NT2RM2001544	0. 59
	NT2RM2001547	F-NT2RM2001547	0. 48
40	NT2RM2001575	F-NT2RM2001575	0. 91
40	NT2RM2001582	F-NT2RM2001582	0. 34
	NT2RM2001588	F-NT2RM2001588	0. 48
	NT2RM2001592	F-NT2RM2001592	0.64
	NT2RM2001605	F-NT2RM2001605	0. 47
45	NT2RM2001613	F-NT2RM2001613	0. 61
	NT2RM2001632	F-NT2RM2001632	0. 31
	NT2RM2001635	F-NT2RM2001635	0. 39
	NT2RM2001637	F-NT2RM2001637	0. 71
50	NT2RM2001641	F-NT2RM2001641	0. 73
	NT2RM2001648	F-NT2RM2001648	0.44
	NT2RM2001652	F-NT2RM2001652	0.40
	NT2RM2001659	F-NT2RM2001659	0. 35
55	NT2RM2001664	F-NT2RM2001664	0. 53
	NT2RM2001668	F-NT2RM2001668	0. 64

	NT2RM2001670	F-NT2RM2001670	0.59 ·
	NT2RM2001671	F-NT2RM2001671	0. 41
	NT2RM2001675	F-NT2RM2001675	0.51
5	NT2RM2001681	F-NT2RM2001681	0. 33
	NT2RM2001688	F-NT2RM2001688	0.74
	NT2RM2001695	F-NT2RM2001695	0. 31
	NT2RM2001696	F-NT2RM2001696	0. 94
10	NT2RM2001698	F-NT2RM2001698	0. 69
	NT2RM2001699	F-NT2RM2001699	0. 85
	NT2RM2001700	F-NT2RM2001700	0.71
	NT2RM2001706	F-NT2RM2001706	0.38
15	NT2RM2001716	F-NT2RM2001716	0.78
	NT2RM2001718	F-NT2RM2001718	0. 81
	NT2RM2001723	F-NT2RM2001723,	0, 83
	NT2RM2001727	F-NT2RM2001727	0. 87
20	NT2RM2001730	F-NT2RM2001730	0. 94
	NT2RM2001743	F-NT2RM2001743	0.94
	NT2RM2001753	F-NT2RM2001753	0. 75
	NT2RM2001760	F-NT2RM2001760	0. 75
25	NT2RM2001768	F-NT2RM2001768	0.71
	NT2RM2001771	F-NT2RM2001771	0. 33
	NT2RM2001782	F-NT2RM2001782	0. 34
	NT2RM2001784	F-NT2RM2001784	0. 45
30	NT2RM2001785	F-NT2RM2001785	0.94
	NT2RM2001797	F-NT2RM2001797	0. 74
	NT2RM2001800	F-NT2RM2001800	0. 90
	NT2RM2001803	F-NT2RM2001803	0. 36
35	NT2RM2001805	F-NT2RM2001805	0. 81
	NT2RM2001813	F-NT2RM2001813	0. 94
	NT2RM2001823	F-NT2RM2001823	0. 94
	NT2RM2001839 NT2RM2001840	F-NT2RM2001839 F-NT2RM2001840	0. 39 0. 94
40	NT2RM2001840	F-NT2RM2001855	0. 94
40	NT2RM2001867	F-NT2RM2001867	0. 92
	NT2RM2001879	F-NT2RM2001879	0. 82
	NT2RM2001886	F-NT2RM2001886	0. 91
45	NT2RM2001896	F-NT2RM2001896	0. 94
45	NT2RM2001903	F-NT2RM2001903	0. 61
	NT2RM2001930	F-NT2RM2001930	0. 94
	NT2RM2001935	F-NT2RM2001935	0.77
50	NT2RM2001936	F-NT2RM2001936	0. 94
50	NT2RM2001950	F-NT2RM2001950	0.34
	NT2RM2001982	F-NT2RM2001982	0. 48
	NT2RM2001983	F-NT2RM2001983	0. 59
	NT2RM2001989	F-NT2RM2001989	0. 94
55	NT2RM2001997	F-NT2RM2001997	0. 90

	NT2RM2001998	F-NT2RM2001998	0. 72
	NT2RM2002004	F-NT2RM2002004	0. 46
5	NT2RM2002014	F-NT2RM2002014	0. 32
3	NT2RM2002030	F-NT2RM2002030_	0. 54
•	NT2RM2002049	F-NT2RM2002049	0. 59
	NT2RM2002055	F-NT2RM2002055	0.94
	NT2RM2002088	F-NT2RM2002088	0. 47
10	NT2RM2002091	F-NT2RM2002091	0. 78
	NT2RM2002100	F-NT2RM2002100	0. 87
	NT2RM2002109	F-NT2RM2002109	0.86
	NT2RM2002128	F-NT2RM2002128	0. 42
15	NT2RM2002142	F-NT2RM2002142	0. 94
	NT2RM2002145	F-NT2RM2002145	0. 94
	NT2RM2002178	F-NT2RM2002178;	0. 44
	NT2RM2002580	F-NT2RM2002580	0. 94
20	NT2RM4000024	F-NT2RM4000024	0. 76
	NT2RM4000027	F-NT2RM4000027	0. 40
	NT2RM4000030	F-NT2RM4000030	0. 66
	NT2RM4000061	F-NT2RM4000061	0. 33
25	NT2RM4000104	F-NT2RM4000104	0. 93
	NT2RM4000139	F-NT2RM4000139	0.87
	NT2RM4000155	F-NT2RM4000155	0.51
	NT2RM4000156	F-NT2RM4000156	0.39
30	NT2RM4000167	F-NT2RM4000167	0. 50
	NT2RM4000169	F-NT2RM4000169	0. 44
	NT2RM4000191	F-NT2RM4000191	0. 94
	NT2RM4000197	F-NT2RM4000197	0.41
35	NT2RM4000199	F-NT2RM4000199	0. 76
	NT2RM4000202	F-NT2RM4000202	0.47
	NT2RM4000210	F-NT2RM4000210	0.44
	NT2RM4000215	F-NT2RM4000215	0.94
40	NT2RM4000229 NT2RM4000233	F-NT2RM4000229 F-NT2RM4000233	0. 93 0. 74
	NT2RM4000251	F-NT2RM4000251	0. 48
	NT2RM4000290	F-NT2RM4000291	0. 46
	NT2RM4000324	F-NT2RM4000324	0. 34
45	NT2RM4000344	F-NT2RM4000344	0. 94
	NT2RM4000349	F-NT2RM4000349	0. 78
	NT2RM4000354	F-NT2RM4000354	0. 76
	NT2RM4000356	F-NT2RM4000356	0.58
50	NT2RM4000386	F-NT2RM4000386	0. 90
	NT2RM4000395	F-NT2RM4000395	0. 47
	NT2RM4000421	F-NT2RM4000421	0.46
	NT2RM4000433	F-NT2RM4000433	0.48
55	NT2RM4000457	F-NT2RM4000457	0. 75
	NT2RM4000471	F-NT2RM4000471	0. 92
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	NT2RM4000486	F-NT2RM4000486	0. 55
	NT2RM4000496	F-NT2RM4000496	0. 62
5	NT2RM4000511	F-NT2RM4000511	0.74
	NT2RM4000514	F-NT2RM4000514_	0.72
	NT2RM4000515	F-NT2RM4000515	0.66
	NT2RM4000520	F-NT2RM4000520	0.69
10	NT2RM4000531	F-NT2RM4000531	0,62
10	NT2RM4000534	F-NT2RM4000534	0.94
	NT2RM4000585	F-NT2RM4000585	0.52
	NT2RM4000590	F-NT2RM4000590	0.94
	NT2RM4000595	F-NT2RM4000595	0.44
15	NT2RM4000603	F-NT2RM4000603	0.94
	NT2RM4000611	F-NT2RM4000611	0.46
	NT2RM4000616	F-NT2RM4000616,	0.94
	NT2RM4000674	F-NT2RM4000674	0.80
20	NT2RM4000689	F-NT2RM4000689	0.58
	NT2RM4000698	F-NT2RM4000698	0.71
	NT2RM4000700	F-NT2RM4000700	0. 59
	NT2RM4000712	F-NT2RM4000712	0.94
25	NT2RM4000717	F-NT2RM4000717	0.69
	NT2RM4000733	F-NT2RM4000733	0. 94
	NT2RM4000734	F-NT2RM4000734	0. 42
	NT2RM4000741	F-NT2RM4000741	0.83
30	NT2RM4000751	F-NT2RM4000751	0.91
	NT2RM4000764	F-NT2RM4000764	0. 35
	NT2RM4000778	F-NT2RM4000778	0.90
	NT2RM4000787	F-NT2RM4000787	0.40
35	NT2RM4000790	F-NT2RM4000790	0.62
	NT2RM4000795	F-NT2RM4000795	0. 94
	NT2RM4000796	F-NT2RM4000796	0. 93
	NT2RM4000798	F-NT2RM4000798	0. 61
40	NT2RM4000813	F-NT2RM4000813	0. 58
40	NT2RM4000820	F-NT2RM4000820	0.39
	NT2RM4000833	F-NT2RM4000833	0.80
	NT2RM4000848	F-NT2RM4000848	0.73
	NT2RM4000852	F-NT2RM4000852	0.74
45	NT2RM4000887	F-NT2RM4000887	0. 42
	NT2RM4000895	F-NT2RM4000895	0. 47
	NT2RM4000950	F-NT2RM4000950	0.53
	NT2RM4000971	F-NT2RM4000971	0.73
50	NT2RM4000996	F-NT2RM4000996	0. 42
	NT2RM4001002	F-NT2RM4001002	0.57
	NT2RM4001032	F-NT2RM4001032	0. 94
	NT2RM4001047	F-NT2RM4001047	0. 76
55	NT2RM4001054	F-NT2RM4001054	0.91
	NT2RM4001084	F-NT2RM4001084	0. 94

	NT2RM4001092	F-NT2RM4001092	0. 94
	NT2RM4001116	F-NT2RM4001116	0.36
	NT2RM4001140	F-NT2RM4001140	0. 82
5	NT2RM4001151	F-NT2RM4001151	0.94
	NT2RM4001155	F-NT2RM4001155	0. 94
	NT2RM4001187	F-NT2RM4001187	0.61
	NT2RM4001191	F-NT2RM4001191	0.77
10	NT2RM4001200	F-NT2RM4001200	0.72
	NT2RM4001203	F-NT2RM4001203	0. 60
	NT2RM4001204	F-NT2RM4001204	0.89
	NT2RM4001217	F-NT2RM4001217	0.93
15	NT2RM4001256	F-NT2RM4001256	0.89
	NT2RM4001258	F-NT2RM4001258	0. 94
	NT2RM4001309	F-NT2RM4001309	
	NT2RM4001316	F-NT2RM4001316	0.38
20	NT2RM4001320	F-NT2RM4001320	0. 33
	NT2RM4001340	F-NT2RM4001340	0.83
	NT2RM4001344	F-NT2RM4001344	0.94
	NT2RM4001347	F-NT2RM4001347	0. 88
25	NT2RM4001371	F-NT2RM4001371	0.61
20	NT2RM4001382	F-NT2RM4001382	0. 93
	NT2RM4001384	F-NT2RM4001384	0.32
	NT2RM4001410	F-NT2RM4001410	0.94
	NT2RM4001411	F-NT2RM4001411	0.47
30	NT2RM4001412	F-NT2RM4001412	0.85
	NT2RM4001444	F-NT2RM4001444	0.94
	NT2RM4001454	F-NT2RM4001454	0.72
	NT2RM4001455	F-NT2RM4001455	0.39
35	NT2RM4001483	F-NT2RM4001483	0. 76
	NT2RM4001489	F-NT2RM4001489	0.94
	NT2RM4001522	F-NT2RM4001522	0.88
	NT2RM4001565	F-NT2RM4001565	0. 78
40	NT2RM4001566	F-NT2RM4001566	0. 87
	NT2RM4001569	F-NT2RM4001569	0.70
	NT2RM4001582	F-NT2RM4001582	0. 76
	NT2RM4001592	F-NT2RM4001592	0. 31
45	NT2RM4001594	F-NT2RM4001594	0.86
	NT2RM4001597	F-NT2RM4001597	0. 91
	NT2RM4001611	F-NT2RM4001611	0. 91
	NT2RM4001629	F-NT2RM4001629	0. 72
50	NT2RM4001650	F-NT2RM4001650	0.37
	NT2RM4001662	F-NT2RM4001662	0.70
	NT2RM4001666	F-NT2RM4001666	0.37
	NT2RM4001682	F-NT2RM4001682	0.91
55	NT2RM4001710	F-NT2RM4001710	0. 45
	NT2RM4001714	F-NT2RM4001714	0. 94

	NT2RM4001715	F-NT2RM4001715	0. 33
	NT2RM4001731	F-NT2RM4001731	0. 94
	NT2RM4001741	F-NT2RM4001741	0. 87
5	NT2RM4001746	F-NT2RM4001746_	
	NT2RM4001758	F-NT2RM4001.758	0. 59
	NT2RM4001783	F-NT2RM4001783	0. 68
	NT2RM4001813	F-NT2RM4001813	0. 39
10	NT2RM4001819	F-NT2RM4001819	0. 43
	NT2RM4001823	F-NT2RM4001823	0. 94
	NT2RM4001828	F-NT2RM4001828	0. 83
	NT2RM4001836	F-NT2RM4001836	0. 61
15	NT2RM4001841	F-NT2RM4001841	0. 37
	NT2RM4001842	F-NT2RM4001842	0. 68
	NT2RM4001865	F-NT2RM4001865	-
	NT2RM4001876	F-NT2RM4001876	0. 88
20	NT2RM4001880	F-NT2RM4001880	0. 43
20	NT2RM4001905	F-NT2RM4001905	0. 94
	NT2RM4001922	F-NT2RM4001922	0. 48
	NT2RM4001938	F-NT2RM4001938	0. 67
	NT2RM4001940	F-NT2RM4001940	0. 92
25	NT2RM4001965	F-NT2RM4001965	0.94
	NT2RM4001969	F-NT2RM4001969	0.88
	NT2RM4001979	F-NT2RM4001979	0.42
	NT2RM4001987	F-NT2RM4001987	0.86
30	NT2RM4002013	F-NT2RM4002013	0. 94
	NT2RM4002018	F-NT2RM4002018	0. 92
	NT2RM4002034	F-NT2RM4002034	0. 43
	NT2RM4002044	F-NT2RM4002044	0.42
35	NT2RM4002054	F-NT2RM4002054	0.88
	NT2RM4002055	F-NT2RM4002055	0.32
	NT2RM4002062	F-NT2RM4002062	0.87
	NT2RM4002063	F-NT2RM4002063	0.49
40	NT2RM4002066	F-NT2RM4002066	0. 39
	NT2RM4002073	F-NT2RM4002073	0. 78
	NT2RM4002075		0.87
	NT2RM4002093	F-NT2RM4002093	0. 82
45	NT2RM4002109	F-NT2RM4002109	0. 52
	NT2RM4002128	F-NT2RM4002128	0. 68
	NT2RM4002140	F-NT2RM4002140	0.94
	NT2RM4002145	F-NT2RM4002145	0.94
50	NT2RM4002146	F-NT2RM4002146	0.89
	NT2RM4002161	F-NT2RM4002161	0.49
	NT2RM4002174	F-NT2RM4002174	0. 36
	NT2RM4002189	F-NT2RM4002189	0. 34
55	NT2RM4002194	F-NT2RM4002194	0. 73
	NT2RM4002205	F-NT2RM4002205	0. 94

	NT2RM4002213	F-NT2RM4002213	0.52
	NT2RM4002226	F-NT2RM4002226	0.48
5	NT2RM4002251	F-NT2RM4002251	0.94
5	NT2RM4002256	F-NT2RM4002256_	_0.93
	NT2RM4002266	F-NT2RM4002266	0.73
	NT2RM4002294	F-NT2RM4002294	0.86
	NT2RM4002301	F-NT2RM4002301	0.40
10	NT2RM4002323	F-NT2RM4002323	0.81
	NT2RM4002339	F-NT2RM4002339	0.32
	NT2RM4002344	F-NT2RM4002344	0.87
	NT2RM4002373	F-NT2RM4002373	0.41
15	NT2RM4002374	F-NT2RM4002374	0.33
	NT2RM4002398	F-NT2RM4002398	0.94
	NT2RM4002409	F-NT2RM4002409	0.82
	NT2RM4002446	F-NT2RM4002446	0. 61
20	NT2RM4002452	F-NT2RM4002452	0.47
	NT2RM4002457	F-NT2RM4002457	0. 92
	NT2RM4002460	F-NT2RM4002460	0.61
	NT2RM4002482	F-NT2RM4002482	0. 94
25	NT2RM4002493	F-NT2RM4002493	0. 39
	NT2RM4002527	F-NT2RM4002527	0. 64
	NT2RM4002534	F-NT2RM4002534	0. 76
	NT2RM4002558	F-NT2RM4002558	0. 62
30	NT2RM4002565	F-NT2RM4002565	0. 54
	NT2RM4002571	F-NT2RM4002571	0.90
	NT2RM4002593	F-NT2RM4002593	0. 91
	NT2RM4002594	F-NT2RM4002594	0.70
35	NT2RM4002623 NT2RP1000018	F-NT2RM4002623	0. 81
	NT2RP1000018	F-NT2RP1000018 F-NT2RP1000035	0.31
	NT2RP1000035	F-NT2RP1000039	0. 49 0. 45
	NT2RP1000040	F-NT2RP1000063	0. 45
40	NT2RP1000086	F-NT2RP1000086	0. 89
	NT2RP1000101	F-NT2RP1000101	0. 76
	NT2RP1000111	F-NT2RP1000111	0. 70
	NT2RP1000112	F-NT2RP1000112	0. 45
45	NT2RP1000124	F-NT2RP1000124	0. 94
	NT2RP1000130	F-NT2RP1000130	0. 69
	NT2RP1000163	F-NT2RP1000163	0. 90
	NT2RP1000170	F-NT2RP1000170	0. 94
50	NT2RP1000174	F-NT2RP1000174	0. 74
	NT2RP1000191	F-NT2RP1000191	0.76
	NT2RP1000202	F-NT2RP1000202	0.60
	NT2RP1000243	F-NT2RP1000243	0. 81
55	NT2RP1000259	F-NT2RP1000259	0.79
	NT2RP1000272	F-NT2RP1000272	0.94

	NT2RP1000324	F-NT2RP1000324	0. 32
	NT2RP1000326	F-NT2RP1000326	0. 94
	NT2RP1000333	F-NT2RP1000333	0. 94
5	NT2RP1000348	F-NT2RP1000348_	0.94
	NT2RP1000357	F-NT2RP1000357	0. 87
	NT2RP1000358	F-NT2RP1000358	0.79
	NT2RP1000363	F-NT2RP1000363	0.40
10	NT2RP1000376	F-NT2RP1000376	0. 61
	NT2RP1000409	F-NT2RP1000409	0. 85
	NT2RP1000413	F-NT2RP1000413	0. 42
	NT2RP1000416	F-NT2RP1000416	0. 68
15	NT2RP1000418	F-NT2RP1000418	0.88
	NT2RP1000439	F-NT2RP1000439	0. 69
	NT2RP1000443	F-NT2RP1000443	0.59
	NT2RP1000460	F-NT2RP1000460	0.86
20	NT2RP1000470	F-NT2RP1000470	0. 56
	NT2RP1000478	F-NT2RP1000478	0.87
	NT2RP1000481	F-NT2RP1000481	0.58
	NT2RP1000493	F-NT2RP1000493	0.34
25	NT2RP1000513	F-NT2RP1000513	0.89
	NT2RP1000522	F-NT2RP1000522	0. 34
	NT2RP1000547	F-NT2RP1000547	0.89
	NT2RP1000574	F-NT2RP1000574	0.74
30	NT2RP1000577	F-NT2RP1000577	0.53
	NT2RP1000581	F-NT2RP1000581	0.66
	NT2RP1000609	F-NT2RP1000609	0.90
	NT2RP1000629	F-NT2RP1000629	0.94
35	NT2RP1000630 NT2RP1000677	F-NT2RP1000630	0.93
	NT2RP1000677	F-NT2RP1000677	0.94
	NT2RP1000685	F-NT2RP1000688 F-NT2RP1000695	0. 94 0. 89
	NT2RP1000701	F-NT2RP1000701	0. 89
40	NT2RP1000721	F-NT2RP1000721	0. 91
	NT2RP1000730	F-NT2RP1000730	0. 94
	NT2RP1000733	F-NT2RP1000733	
	NT2RP1000738	F-NT2RP1000738	0. 64
45	NT2RP1000746		0. 92
	NT2RP1000767	F-NT2RP1000767	0. 82
	NT2RP1000782	F-NT2RP1000782	0. 94
	NT2RP1000796	F-NT2RP1000796	0. 33
50	NT2RP1000825	F-NT2RP1000825	0.74
••	NT2RP1000833	F-NT2RP1000833	0. 94
	NT2RP1000834	F-NT2RP1000834	0. 88
	NT2RP1000836	F-NT2RP1000836	0. 59
55	NT2RP1000846	F-NT2RP1000846	0.46
55	NT2RP1000851	F-NT2RP1000851	0.82

	NT2RP1000856	F-NT2RP1000856	0. 94
	NT2RP1000860	F-NT2RP1000860	0.60
5	NT2RP1000902	F-NT2RP1000902	0. 42
	NT2RP1000915	F-NT2RP1000915_	0, 90
	NT2RP1000916	F-NT2RP1000916	0. 67
	NT2RP1000943	F-NT2RP1000943	0.77
	NT2RP1000944	F-NT2RP1000944	0.94
10	NT2RP1000947	F-NT2RP1000947	0. 38
	NT2RP1000954	F-NT2RP1000954	0. 59
	NT2RP1000958	F-NT2RP1000958	0. 59
	NT2RP1000959	F-NT2RP1000959	0.74
15	NT2RP1000966	F-NT2RP1000966	0. 93
	NT2RP1000980	F-NT2RP1000980	0, 31
	NT2RP1000988	F-NT2RP1000988	
	NT2RP1001011	F-NT2RP1001011	0. 65
20	NT2RP1001013	F-NT2RP1001013	0. 34
	NT2RP1001014	F-NT2RP1001014	0. 94
	NT2RP1001033	F-NT2RP1001033	0.40
	NT2RP1001073	F-NT2RP1001073	0.74
25	NT2RP1001079	F-NT2RP1001079	0.48
	NT2RP1001080	F-NT2RP1001080	0. 91
	NT2RP1001113	F-NT2RP1001113	0. 94
	NT2RP1001173	F-NT2RP1001173	0.57
30	NT2RP1001177	F-NT2RP1001177	0.75
	NT2RP1001185	F-NT2RP1001185	0. 94
	NT2RP1001199	F-NT2RP1001199	0. 42
	NT2RP1001247	F-NT2RP1001247	0. 67
35	NT2RP1001248	F-NT2RP1001248	0. 89
	NT2RP1001253	F-NT2RP1001253	0. 94
	NT2RP1001286	F-NT2RP1001286	0. 59
	NT2RP1001294	F-NT2RP1001294	0. 81
40	NT2RP1001302	F-NT2RP1001302	0. 86
40	NT2RP1001310	F-NT2RP1001310	0. 91
	NT2RP1001311	F-NT2RP1001311	0. 87
	NT2RP1001313	F-NT2RP1001313	0. 91
	NT2RP1001361	F-NT2RP1001361	0.83
45	NT2RP1001385	F-NT2RP1001385	0. 36
	NT2RP1001395	F-NT2RP1001395	0.84
	NT2RP1001410	F-NT2RP1001410	0.72
	NT2RP1001424	F-NT2RP1001424	0.81
50	NT2RP1001432	F-NT2RP1001432	0.81
	NT2RP1001449	F-NT2RP1001449	0.77
	NT2RP1001457	F-NT2RP1001457	0. 92
	NT2RP1001466	F-NT2RP1001466	0.44
55	NT2RP1001475	F-NT2RP1001475	0.53
	NT2RP1001482	F-NT2RP1001482	0. 34

	NT2RP1001494	F-NF2RP1001494	0. 93
	NT2RP1001543	F-NT2RP1001543	0. 94
5	NT2RP1001546	F-NT2RP1001546	0.94
	NT2RP1001569	F-NT2RP1001569_	0, 93
	NT2RP1001616	F-NT2RP1001616	0. 46
	NT2RP1001665	F-NT2RP1001665	0. 94
	NT2RP2000001	F-NT2RP2000001	0. 94
10	NT2RP2000006	F-NT2RP2000006	0. 74
	NT2RP2000007	F-NT2RP2000007	0. 94
	NT2RP2000008	F-NT2RP2000008	0. 83
	NT2RP2000032	F-NT2RP2000032	0. 94
15	NT2RP2000040	F-NT2RP2000040	0. 92
	NT2RP2000045	F~NT2RP2000045	0. 94
	NT2RP2000054	F-NT2RP2000054,	0. 53
	NT2RP2000056	F-NT2RP2000056	0. 42
20	NT2RP2000067	F-NT2RP2000067	0. 88
	NT2RP2000070	F-NT2RP2000070	0. 48
	NT2RP2000079	F-NT2RP2000079	0. 85
	NT2RP2000088	F-NT2RP2000088	0. 94
25	NT2RP2000091	F-NT2RP2000091	0. 79
	NT2RP2000097	F-NT2RP2000097	0.66
	NT2RP2000114	F-NT2RP2000114	0.86
	NT2RP2000120	F-NT2RP2000120	0. 48
30	NT2RP2000126	F-NT2RP2000126	0. 77
	NT2RP2000133	F-NT2RP2000133	0. 64
	NT2RP2000147	F-NT2RP2000147	0. 94
	NT2RP2000153	F-NT2RP2000153	0. 87
35	NT2RP2000157	F-NT2RP2000157	0. 48
	NT2RP2000161	F-NT2RP2000161	0.79
	NT2RP2000173 NT2RP2000175	F-NT2RP2000173	0. 75
	NT2RP2000175	F-NT2RP2000175	0.79
40	NT2RP2000195	F-NT2RP2000195 F-NT2RP2000205	0.91
	NT2RP2000208	F-NT2RP2000208	0. 94 0. 84
	NT2RP2000208	F-NT2RP2000208	0. 52
	NT2RP2000232	F-NT2RP2000232	0. 32
45	NT2RP2000233	F-NT2RP2000233	0. 57
	NT2RP2000239	F-NT2RP2000239	0. 52
	NT2RP2000248	F-NT2RP2000248	0. 36
	NT2RP2000274	F-NT2RP2000274	0. 55
50	NT2RP2000283	F-NT2RP2000283	0. 61
	NT2RP2000288	F-NT2RP2000288	0. 31
	NT2RP2000297	F-NT2RP2000297	0. 78
	NT2RP2000298	F-NT2RP2000298	0. 94
55	NT2RP2000310	F-NT2RP2000310	0. 52
	NT2RP2000328	F-NT2RP2000328	0.75

	NT2RP2000329	F-NT2RP2000329	0.94
	NT2RP2000346	F-NT2RP2000346	0. 62
_	NT2RP2000369	F-NT2RP2000369	0.87
5	NT2RP2000412	F-NT2RP2000412	0. 75
	NT2RP2000414	F-NT2RP2000414	0. 79
	NT2RP2000422	F-NT2RP2000422	0. 46
	NT2RP2000438	F-NT2RP2000438	0. 94
10	NT2RP2000448	F-NT2RP2000448	0.94
	NT2RP2000503	F-NT2RP2000503	0. 94
	NT2RP2000510	F-NT2RP2000510	0. 87
	NT2RP2000516	F-NT2RP2000516	0. 42
15	NT2RP2000617	F-NT2RP2000617	0.94
	NT2RP2000634	F-NT2RP2000634	0.74
	NT2RP2000656	F-NT2RP2000656,	0. 68
	NT2RP2000658	F-NT2RP2000658	0. 55
20	NT2RP2000668	F-NT2RP2000668	0. 81
	NT2RP2000704	F-NT2RP2000704	0. 88
	NT2RP2000710	F-NT2RP2000710	0. 58
	NT2RP2000764	F-NT2RP2000764	0. 59
25	NT2RP2000809	F-NT2RP2000809	0.51
	NT2RP2000812	F-NT2RP2000812	0. 37
	NT2RP2000814	F-NT2RP2000814	0.66
	NT2RP2000816	F-NT2RP2000816	0.50
30	NT2RP2000819	F-NT2RP2000819	0. 35
30	NT2RP2000841	F-NT2RP2000841	0. 93
	NT2RP2000845	F-NT2RP2000845	0. 78
	NT2RP2000863	F-NT2RP2000863	0. 41
	NT2RP2000880	F-NT2RP2000880	0. 79
35	NT2RP2000892	F-NT2RP2000892	0. 94
	NT2RP2000931	F-NT2RP2000931	0. 89
	NT2RP2000932	F-NT2RP2000932	0. 94
	NT2RP2000938	F-NT2RP2000938	0. 90
40	NT2RP2000943	F-NT2RP2000943	0. 61
	NT2RP2000965	F-NT2RP2000965	0. 80
	NT2RP2000985	F-NT2RP2000985	0. 94
	NT2RP2001036	F-NT2RP2001036	0. 86
45	NT2RP2001044	F-NT2RP2001044	0. 93
	NT2RP2001056	F-NT2RP2001056	0. 77
	NT2RP2001065	F-NT2RP2001065	0. 73
	NT2RP2001070	F-NT2RP2001070	0. 91
50	NT2RP2001081	F-NT2RP2001081	0. 94
	NT2RP2001094	F-NT2RP2001094	0. 93
	NT2RP2001119	F-NT2RP2001119	0. 31
	NT2RP2001127	F-NT2RP2001127	0. 41
55	NT2RP2001137	F-NT2RP2001137	0.93
	NT2RP2001168	F-NT2RP2001168	0. 71

	NT2RP2001173	F-NT2RP2001173	0. 94
	NT2RP2001174	F-NT2RP2001174	0.71
	NT2RP2001218	F-NT2RP2001218	0.76
5	NT2RP2001233		
	NT2RP2001245	F-NT2RP2001245	_
	NT2RP2001268	F-NT2RP2001268	0. 61
	NT2RP2001327	F-NT2RP2001327	0. 86
10	NT2RP2001328	F-NT2RP2001328	0. 94
	NT2RP2001366	F-NT2RP2001366	0. 52
	NT2RP2001378	F-NT2RP2001378	0. 64
	NT2RP2001381	F-NT2RP2001381	0.80
15	NT2RP2001392	F-NT2RP2001392	0.69
	NT2RP2001394	F-NT2RP2001394	0. 38
	NT2RP2001397	F-NT2RP2001397,	
	NT2RP2001420	F-NT2RP2001420	0.88
20	NT2RP2001427	F-NT2RP2001427	0. 73
	NT2RP2001436	F-NT2RP2001436	0. 32
	NT2RP2001440	F-NT2RP2001440	0. 94
	NT2RP2001450	F-NT2RP2001450	0.38
25	NT2RP2001511	F-NT2RP2001511	0. 53
	NT2RP2001520	F-NT2RP2001520	0.88
	NT2RP2001536	F-NT2RP2001536	0.64
	NT2RP2001560	F-NT2RP2001560	0.79
30	NT2RP2001576	F-NT2RP2001576	0.62
30	NT2RP2001581	F-NT2RP2001581	0.48
	NT2RP2001597	F-NT2RP2001597	0.94
	NT2RP2001601	F-NT2RP2001601	0. 33
0.5	NT2RP2001613	F-NT2RP2001613	0. 41
35	NT2RP2001628	F-NT2RP2001628	0. 79
	NT2RP2001634	F-NT2RP2001634	0. 87
	NT2RP2001660	F-NT2RP2001660	0.86
	NT2RP2001675	F-NT2RP2001675	0. 90
40	NT2RP2001721	F-NT2RP2001721	0.32
	NT2RP2001748	F-NT2RP2001748	0. 32
	NT2RP2001839	1 1112111 200 1000	0. 83
	NT2RP2001861		0. 33
45	NT2RP2001876	F-NT2RP2001876	0. 94
	NT2RP2001883	F-NT2RP2001883	0. 90
	NT2RP2001898	F-NT2RP2001898	0. 94
	NT2RP2001900	F-NT2RP2001900	0. 93
50	NT2RP2001907	F-NT2RP2001907	0. 94
	NT2RP2001946	F-NT2RP2001946	0. 63
	NT2RP2001947	F-NT2RP2001947	0. 75
	NT2RP2001969	F-NT2RP2001969	0.94
55	NT2RP2001976	F-NT2RP2001976	0. 48
	NT2RP2001985	F-NT2RP2001985	0. 44

	NT2RP2001991	F-NT2RP2001991	0. 39
	NT2RP2002025	F-NT2RP2002025	0.94
_	NT2RP2002033	F-NT2RP2002033	0.40
5	NT2RP2002046	F-NT2RP2002046_	0. 56
	NT2RP2002058	F-NT2RP2002058	0.64
•	NT2RP2002076	F-NT2RP2002076	0.40
	NT2RP2002078	F-NT2RP2002078	0. 81
10	NT2RP2002079	F-NT2RP2002079	0. 82
	NT2RP2002099	F-NT2RP2002099	0. 92
	NT2RP2002154	F-NT2RP2002154	0. 44
	NT2RP2002185	F-NT2RP2002185	0.94
15	NT2RP2002193	F~NT2RP2002193	0. 48
	NT2RP2002208	F-NT2RP2002208	0. 49
	NT2RP2002231	F-NT2RP2002231	0. 61
	NT2RP2002235	F-NT2RP2002235	0. 87
20	NT2RP2002252	F-NT2RP2002252	0.51
	NT2RP2002270	F-NT2RP2002270	0.78
	NT2RP2002292	F-NT2RP2002292	0.94
	NT2RP2002312	F-NT2RP2002312	0.33
25	NT2RP2002325	F-NT2RP2002325	0. 53
	NT2RP2002333	F-NT2RP2002333	0.60
	NT2RP2002373	F-NT2RP2002373	0.91
	NT2RP2002385	F-NT2RP2002385	0.94
30	NT2RP2002408	F-NT2RP2002408	0.44
	NT2RP2002426	F-NT2RP2002426	0. 54
	NT2RP2002442	F-NT2RP2002442	0. 37
	NT2RP2002464	F-NT2RP2002464	0. 79
35	NT2RP2002479	F-NT2RP2002479	0. 92
	NT2RP2002498	F-NT2RP2002498	0.94
	NT2RP2002503	F-NT2RP2002503	0. 90
	NT2RP2002520	F-NT2RP2002520	0.37
40	NT2RP2002537	F-NT2RP2002537	0. 42
	NT2RP2002549 NT2RP2002595	F-NT2RP2002549	0.75
	NT2RP2002609	F-NT2RP2002595	0.90
	NT2RP2002609	F-NT2RP2002609 F-NT2RP2002618	
45	NT2RP2002621	F-NT2RP2002618	0. 52 0. 67
•	NT2RP2002672	F-NT2RP2002672	0. 43
	NT2RP2002701	F-NT2RP2002701	0. 43
	NT2RP2002706	F-NT2RP2002706	0. 51
50	NT2RP2002710	F-NT2RP2002710	0. 94
	NT2RP2002769	F-NT2RP2002769	0. 42
	NT2RP2002800	F-NT2RP2002800	0. 47
	NT2RP2002862	F-NT2RP2002862	0. 47
	NT2RP2002880	F-NT2RP2002880	0. 78
55	NT2RP2002891	F-NT2RP2002891	0. 69
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	NT2RP2002925	F-NT2RP2002925	0. 65
	NT2RP2002928		0. 47
	NT2RP2002929	F-NT2RP2002928 F-NT2RP2002929	0. 47
5			
	NT2RP2002939		•
	NT2RP2002954	F-NT2RP2002954	0.94
	NT2RP2002959	F-NT2RP2002959	0. 45
10	NT2RP2002979	F-NT2RP2002979	0. 51
10	NT2RP2002980	F-NT2RP2002980	0.38
	NT2RP2002986		0.34
	NT2RP2002993	F-NT2RP2002993	0. 94
	NT2RP2003034	F-NT2RP2003034	0. 34
15	NT2RP2003099	F-NT2RP2003099	0. 46
	NT2RP2003108		0. 67
	NT2RP2003117	F-NT2RP2003117,	0.57
	NT2RP2003121	F-NT2RP2003121	0.94
20	NT2RP2003125	F-NT2RP2003125	0.94
	NT2RP2003137	F-NT2RP2003137	0.60
	NT2RP2003157	F-NT2RP2003157	0.94
	NT2RP2003158	F-NT2RP2003158	0.75
25	NT2RP2003165	F-NT2RP2003165	0.90
	NT2RP2003177	F-NT2RP2003177	0. 85
	NT2RP2003194	F-NT2RP2003194	0.94
	NT2RP2003228	F-NT2RP2003228	0. 57
20	NT2RP2003243	F-NT2RP2003243	0.94
30	NT2RP2003265	F-NT2RP2003265	0.68
	NT2RP2003272	F-NT2RP2003272	0.94
	NT2RP2003277	F-NT2RP2003277	0.73
	NT2RP2003280	F-NT2RP2003280	0.94
35	NT2RP2003286	F-NT2RP2003286	0.44
	NT2RP2003293	F-NT2RP2003293	0.42
	NT2RP2003295	F-NT2RP2003295	0.89
	NT2RP2003297	F-NT2RP2003297	0.94
40	NT2RP2003307	F-NT2RP2003307	0. 79
	NT2RP2003308	F-NT2RP2003308	0.49
	NT2RP2003329	F-NT2RP2003329	0.33
	NT2RP2003347	F-NT2RP2003347	0.79
45	NT2RP2003367	F-NT2RP2003367	0.94
	NT2RP2003391	F-NT2RP2003391	0.90
	NT2RP2003393	F-NT2RP2003393	0.94
	NT2RP2003401	F-NT2RP2003401	0.80
50	NT2RP2003433	F-NT2RP2003433	0.40
	NT2RP2003445	F-NT2RP2003445	0.94
	NT2RP2003446	F-NT2RP2003446	0.40
	NT2RP2003466	F-NT2RP2003466	0.58
==	NT2RP2003480	F-NT2RP2003480	0. 59
55	NT2RP2003506	F-NT2RP2003506	0.94

	NT2RP2003511	F-NT2RP2003511	0: 31
	NT2RP2003513	F-NT2RP2003513	0. 67
5	NT2RP2003533	F-NT2RP2003533	0. 57
	NT2RP2003543	F-NT2RP2003543	0.94
	NT2RP2003564	F-NT2RP2003564	0.88
	NT2RP2003567	F-NT2RP2003567	0.69
	NT2RP2003581	F-NT2RP2003581	0.40
10	NT2RP2003596	F-NT2RP2003596	0.83
	NT2RP2003604	F-NT2RP2003604	0.87
	NT2RP2003629	F-NT2RP2003629	0.87
	NT2RP2003643	F-NT2RP2003643	0.94
15	NT2RP2003687	F-NT2RP2003687	0.31
	NT2RP2003691	F-NT2RP2003691	0.34
	NT2RP2003702	F-NT2RP2003702,	0.80
	NT2RP2003713	F-NT2RP2003713	0. 53
20	NT2RP2003714	F-NT2RP2003714	0. 61
	NT2RP2003737	F-NT2RP2003737	0.88
	NT2RP2003751	F-NT2RP2003751	0. 36
	NT2RP2003760	F-NT2RP2003760	0. 62
25	NT2RP2003764	F-NT2RP2003764	0. 94
	NT2RP2003769	F-NT2RP2003769	0. 69
	NT2RP2003777	F-NT2RP2003777	0. 93
	NT2RP2003781	F-NT2RP2003781	0. 37
30	NT2RP2003793	F-NT2RP2003793	0. 92
	NT2RP2003825	F-NT2RP2003825	0.49
	NT2RP2003840 NT2RP2003857	F-NT2RP2003840	0.88
	NT2RP2003952	F-NT2RP2003857 F-NT2RP2003952	0. 34 0. 89
35	NT2RP2003932	F-NT2RP2003976	0. 89
	NT2RP2003981	F-NT2RP2003981	0. 64
	NT2RP2003984	F-NT2RP2003984	0. 41
	NT2RP2003986	F-NT2RP2003986	0. 32
40	NT2RP2004013	F-NT2RP2004013	0. 48
	NT2RP2004041	F-NT2RP2004041	0. 78
	NT2RP2004042	F-NT2RP2004042	0, 66
	NT2RP2004066	F-NT2RP2004066	0. 56
45	NT2RP2004081	F-NT2RP2004081	0.94
	NT2RP2004098	F-NT2RP2004098	0.39
	NT2RP2004124	F-NT2RP2004124	0.75
	NT2RP2004152	F-NT2RP2004152	0.54
50	NT2RP2004165	F-NT2RP2004165	0.68
	NT2RP2004187	F-NT2RP2004187	0.92
	NT2RP2004194	F-NT2RP2004194	0.90
	NT2RP2004196	F-NT2RP2004196	0. 94
55	NT2RP2004226	F-NT2RP2004226	0. 80
	NT2RP2004239	F-NT2RP2004239	0. 63

	NT2RP2004240	F-NT2RP2004240	0.86
	NT2RP2004242	F-NT2RP2004242	0. 94
_	NT2RP2004245	F-NT2RP2004245	0. 92
5	NT2RP2004316	F-NT2RP2004316_	0. 33
	NT2RP2004364	F-NT2RP2004364	0. 75
	NT2RP2004365	F-NT2RP2004365	0. 43
	NT2RP2004366	F-NT2RP2004366	0. 32
10	NT2RP2004373	F-NT2RP2004373	0. 60
	NT2RP2004389	F-NT2RP2004389	0.94
	NT2RP2004392	F-NT2RP2004392	0. 93
	NT2RP2004399	F-NT2RP2004399	0.36
15	NT2RP2004463	F-NT2RP2004463	0. 43
	NT2RP2004476	F-NT2RP2004476	0. 80
	NT2RP2004538	F-NT2RP2004538	
	NT2RP2004551	F-NT2RP2004551	0. 47
20	NT2RP2004568	F-NT2RP2004568	0. 73
	NT2RP2004594	F-NT2RP2004594	0. 64
	NT2RP2004600	F-NT2RP2004600	0. 53
	NT2RP2004602	F-NT2RP2004602	0. 33
25	NT2RP2004614		0. 34
	NT2RP2004655	F-NT2RP2004655	0. 77
	NT2RP2004664	F-NT2RP2004664	0. 94
	NT2RP2004689	F-NT2RP2004689	0. 83
30	NT2RP2004710	F-NT2RP2004710	0. 94
30	NT2RP2004743	F-NT2RP2004743	0. 32
	NT2RP2004768	F-NT2RP2004768	0. 53
	NT2RP2004791	F-NT2RP2004791	0. 64
	NT2RP2004799	F-NT2RP2004799	0. 42
35	NT2RP2004802	F-NT2RP2004802	0. 88
	NT2RP2004816	F-NT2RP2004816	0. 78
	NT2RP2004841	F-NT2RP2004841	0.61
	NT2RP2004861	F-NT2RP2004861	0. 32
40	NT2RP2004897	F-NT2RP2004897	0. 67
	NT2RP2004933	F-NT2RP2004933	0. 94
	NT2RP2004936	F-NT2RP2004936	0. 53
	NT2RP2004959	F-NT2RP2004959	0. 78
45	NT2RP2004978	F-NT2RP2004978	0. 86
	NT2RP2004985	F-NT2RP2004985	0. 42
	NT2RP2004999	F-NT2RP2004999	0. 55
	NT2RP2005000	F-NT2RP2005000	0.40
50	NT2RP2005001	F-NT2RP2005001	0. 94
	NT2RP2005003	F-NT2RP2005003	0. 44
	NT2RP2005012	F-NT2RP2005012	0. 73
	NT2RP2005018	F-NT2RP2005018	0.43
55	NT2RP2005022	F-NT2RP2005022	0.34
	NT2RP2005037	F-NT2RP2005037	0. 94

	NT2RP2005038	F-NT2RP2005038	0. 56
	NT2RP2005116	F-NT2RP2005116	0. 94
	NT2RP2005126	F-NT2RP2005126	0. 69
5	NT2RP2005139	F-NT2RP2005139	0. 94
	NT2RP2005140	F-NT2RP2005140	0. 32
	NT2RP2005144	F-NT2RP2005144	0. 39
	NT2RP2005147	F-NT2RP2005147	0. 32
10	NT2RP2005159	F-NT2RP2005159	0. 52
	NT2RP2005162	F-NT2RP2005162	0. 86
	NT2RP2005168	F-NT2RP2005168	0. 53
	NT2RP2005204	F-NT2RP2005204	0.94
15	NT2RP2005227	F-NT2RP2005227	0. 53
	NT2RP2005239	F-NT2RP2005239	0. 91
	NT2RP2005270	F-NT2RP2005270,	0.72
	NT2RP2005276	F-NT2RP2005276	0.41
20	NT2RP2005287	F-NT2RP2005287	0.44
-	NT2RP2005288	F-NT2RP2005288	0.82
	NT2RP2005315	F-NT2RP2005315	0.79
	NT2RP2005358	F-NT2RP2005358	0.94
25	NT2RP2005393	F-NT2RP2005393	0.41
20	NT2RP2005436	F-NT2RP2005436	0. 45
	NT2RP2005441	F-NT2RP2005441	0.88
	NT2RP2005453	F-NT2RP2005453	0. 59
30	NT2RP2005457	F-NT2RP2005457	0. 56
30	NT2RP2005464	F-NT2RP2005464	0. 72
	NT2RP2005465	F-NT2RP2005465	0. 40
	NT2RP2005472	F-NT2RP2005472	0.42
	NT2RP2005490	F-NT2RP2005490	0. 52
35	NT2RP2005495	F-NT2RP2005495	0. 94
	NT2RP2005496	F-NT2RP2005496	0.52
	NT2RP2005498	F-NT2RP2005498	0.74
	NT2RP2005509	F-NT2RP2005509	0.55
40	NT2RP2005520	F-NT2RP2005520	0.54
	NT2RP2005525	F-NT2RP2005525	0.41
	NT2RP2005539	F-NT2RP2005539	0.94
	NT2RP2005540	F-NT2RP2005540	0. 32
45	NT2RP2005549 NT2RP2005555	F-NT2RP2005549 F-NT2RP2005555	0. 79
	NT2RP2005557	F-NT2RP2005557	0. 63
	NT2RP2005600	F-NT2RP2005600	0. 76
	NT2RP2005605	F-NT2RP2005605	0. 76
50	NT2RP2005620	F-NT2RP2005620	0.66
	NT2RP2005622	F-NT2RP2005622	0. 37
	NT2RP2005635	F-NT2RP2005635	0. 37
	NT2RP2005637	F-NT2RP2005637	0. 94
55	NT2RP2005640	F-NT2RP2005640	0. 87

	NT2RP2005654	F-NT2RP2005654	0. 94
	NT2RP2005669	F-NT2RP2005669	0.94
	NT2RP2005675	F-NT2RP2005675	0. 72
5	NT2RP2005683	F-NT2RP2005683	0. 72
	NT2RP2005690		•
		F-NT2RP2005690	0.54
	NT2RP2005712	F-NT2RP2005712	0.93
10	NT2RP2005722	F-NT2RP2005722	0.50
10	NT2RP2005723	F-NT2RP2005723	0. 45
	NT2RP2005732	F-NT2RP2005732	0. 54
	NT2RP2005748	F-NT2RP2005748	0. 76
	NT2RP2005752	F-NT2RP2005752	0. 47
15	NT2RP2005763	F-NT2RP2005763	0. 87
	NT2RP2005767		0. 58
	NT2RP2005773	F-NT2RP2005773	0. 49
	NT2RP2005775	F-NT2RP2005775	0.34
20	NT2RP2005781	F-NT2RP2005781	0. 81
	NT2RP2005784	F-NT2RP2005784	0.94
	NT2RP2005804	F-NT2RP2005804	0.94
	NT2RP2005812	F-NT2RP2005812	0.90
25	NT2RP2005835	F-NT2RP2005835	0.94
	NT2RP2005853	F-NT2RP2005853	0.77
	NT2RP2005859	F-NT2RP2005859	0.68
	NT2RP2005868	F-NT2RP2005868	0.76
	NT2RP2005886	F-NT2RP2005886	0.62
30	NT2RP2005890	F-NT2RP2005890	0. 79
	NT2RP2005901	F-NT2RP2005901	0. 94
	NT2RP2005933	F-NT2RP2005933	0. 55
	NT2RP2006023	F-NT2RP2006023	0. 47
35	NT2RP2006038	F-NT2RP2006038	0.73
	NT2RP2006043	F-NT2RP2006043	0.50
	NT2RP2006052	F-NT2RP2006052	0.74
	NT2RP2006069	F-NT2RP2006069	0. 85
40	NT2RP2006071	F-NT2RP2006071	0.94
	NT2RP2006100	F-NT2RP2006100	0.42
	NT2RP2006106	F-NT2RP2006106	0.94
	NT2RP2006141	F-NT2RP2006141	0.92
45	NT2RP2006186	F-NT2RP2006186	0.94
	NT2RP2006196	F-NT2RP2006196	0.32
	NT2RP2006200	F-NT2RP2006200	0.70
	NT2RP2006219	F-NT2RP2006219	0.94
50	NT2RP2006237	F-NT2RP2006237	0. 59
50	NT2RP2006238	F-NT2RP2006238	0. 47
	NT2RP2006275	F-NT2RP2006275	0.89
	NT2RP2006312	F-NT2RP2006312	0. 75
	NT2RP2006333	F-NT2RP2006333	0. 73
55	NT2RP2006334	F-NT2RP2006334	0. 72
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·	NT2RP2006365	F-NT2RP2006365	0. 48
	NT2RP2006436	F-NT2RP2006436	0. 75
	NT2RP2006441	F-NT2RP2006441	0. 61
5	NT2RP2006456	F-NT2RP2006456	
	NT2RP2006464	F-NT2RP2006464	0. 51
	NT2RP2006472	F-NT2RP2006472	0. 76
	NT2RP2006565	F-NT2RP2006565	0. 34
10	NT2RP2006571	F-NT2RP2006571	0. 94
	NT2RP2006573	F-NT2RP2006573	0. 35
	NT2RP3000002	F-NT2RP3000002	0. 40
	NT2RP3000031	F-NT2RP3000031	0. 49
15	NT2RP3000046	F-NT2RP3000046	0. 90
	NT2RP3000047	F-NT2RP3000047	0. 69
	NT2RP3000050	F-NT2RP3000050,	
	NT2RP3000068	F-NT2RP3000068	0. 94
20	NT2RP3000072	F-NT2RP3000072	0. 34
	NT2RP3000080	F-NT2RP3000080	0. 32
	NT2RP3000085	F-NT2RP3000085	0. 92
	NT2RP3000092	F-NT2RP3000092	0. 40
25	NT2RP3000134	F-NT2RP3000134	0.42
	NT2RP3000142	F-NT2RP3000142	0. 58
	NT2RP3000149	F-NT2RP3000149	0.83
	NT2RP3000197	F-NT2RP3000197	0.59
30	NT2RP3000207	F-NT2RP3000207	0.48
30	NT2RP3000220	F-NT2RP3000220	0.75
	NT2RP3000251	F-NT2RP3000251	0.82
	NT2RP3000252	F-NT2RP3000252	0.88
	NT2RP3000255	F-NT2RP3000255	0.94
35	NT2RP3000267	F-NT2RP3000267	0. 92
	NT2RP3000299	F-NT2RP3000299	0.94
	NT2RP3000312	F-NT2RP3000312	0. 57
	NT2RP3000320	F-NT2RP3000320	0. 94
40	NT2RP3000324	F-NT2RP3000324	0.32
	NT2RP3000333	F-NT2RP3000333	0.94
	NT2RP3000348	F-NT2RP3000348	0. 32
	NT2RP3000350	F-NT2RP3000350	0.67
45	NT2RP3000359	F-NT2RP3000359	0.94
	NT2RP3000361	F-NT2RP3000361	0.94
	NT2RP3000366	F-NT2RP3000366	0.94
	NT2RP3000393	F-NT2RP3000393	0. 53
50	NT2RP3000397	F-NT2RP3000397	0.89
	NT2RP3000403	F-NT2RP3000403	0.87
	NT2RP3000439	F-NT2RP3000439	0.77
	NT2RP3000456	F-NT2RP3000456	0.88
55	NT2RP3000484	F-NT2RP3000484	0.59
	NT2RP3000487	F-NT2RP3000487	0. 32

	NT2RP3000526	F-NT2RP3000526	0. 32
	NT2RP3000527	F-NT2RP3000527	0. 69
	NT2RP3000531	F-NT2RP3000531	0. 56
5	NT2RP3000562	F-NT2RP3000562	
	NT2RP3000590	F-NT2RP3000590	0.88
	NT2RP3000592		
		F-NT2RP3000592	0. 92
10	NT2RP3000596	F-NT2RP3000596	0.50
	NT2RP3000599	F-NT2RP3000599	0.69
	NT2RP3000603	F-NT2RP3000603	0. 94
	NT2RP3000605	F-NT2RP3000605	0. 92
	NT2RP3000624	F-NT2RP3000624	0.71
15	NT2RP3000632	F-NT2RP3000632	0. 43
	NT2RP3000644	F-NT2RP3000644	0. 94
	NT2RP3000661	F-NT2RP3000661,	0.60
	NT2RP3000665	F-NT2RP3000665	0. 46
20	NT2RP3000690	F-NT2RP3000690	0. 79
	NT2RP3000739	F-NT2RP3000739	0. 94
	NT2RP3000753	F-NT2RP3000753	0. 42
	NT2RP3000759	F-NT2RP3000759	0. 94
25	NT2RP3000825	F-NT2RP3000825	0. 73
	NT2RP3000826	F-NT2RP3000826	0. 94
	NT2RP3000836	F-NT2RP3000836	0. 39
	NT2RP3000841	F-NT2RP3000841	0. 55
30	NT2RP3000845	F-NT2RP3000845	0. 38
	NT2RP3000847	F-NT2RP3000847	0. 67
	NT2RP3000850	F-NT2RP3000850	0. 59
	NT2RP3000852	F-NT2RP3000852	0. 39
25	NT2RP3000859	F-NT2RP3000859	0. 83
35	NT2RP3000868	F-NT2RP3000868	0. 49
	NT2RP3000869	F-NT2RP3000869	0. 94
	NT2RP3000901	F-NT2RP3000901	0. 34
	NT2RP3000917	F-NT2RP3000917	0.73
40	NT2RP3000919	F-NT2RP3000919	0. 94
	NT2RP3000968	F-NT2RP3000968	0. 94
	NT2RP3000980	F-NT2RP3000980	0. 61
	NT2RP3000994	F-NT2RP3000994	0. 45
45	NT2RP3001004	F-NT2RP3001004	0.86
	NT2RP3001055	F-NT2RP3001055	0. 38
	NT2RP3001057	F-NT2RP3001057	0. 84
	NT2RP3001081	F-NT2RP3001081	0.57
50	NT2RP3001084	F-NT2RP3001084	0.94
	NT2RP3001096	F-NT2RP3001096	0.94
	NT2RP3001107	F-NT2RP3001107	0.88
	NT2RP3001109	F-NT2RP3001109	0.67
55	NT2RP3001111	F-NT2RP3001111	0.87
	NT2RP3001113	F-NT2RP3001113	0.40

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	NT2RP3001116		0. 73
	NT2RP3001119	F-NT2RP3001119	0.62
5	NT2RP3001120	F-NT2RP3001120	0.59
	NT2RP3001126	F-NT2RP3001126	
	NT2RP3001133	F-NT2RP3001133	0.89
	NT2RP3001140	F-NT2RP3001140	0. 59
	NT2RP3001147	F-NT2RP3001147	0.94
10	NT2RP3001150	F-NT2RP3001150	0. 92
	NT2RP3001155	F-NT2RP3001155	0. 94
	NT2RP3001176	F-NT2RP3001176	0. 82
	NT2RP3001214	F-NT2RP3001214	0. 41
15	NT2RP3001216	F-NT2RP3001216	0.32
	NT2RP3001221	F-NT2RP3001221	0.46
	NT2RP3001236	F-NT2RP3001236,	0.79
	NT2RP3001239	F-NT2RP3001239	0.94
20	NT2RP3001253	F-NT2RP3001253	0. 59
	NT2RP3001260	F-NT2RP3001260	0. 94
	NT2RP3001268	F-NT2RP3001268	0.75
	NT2RP3001272	F-NT2RP3001272	0. 94
25	NT2RP3001274	F-NT2RP3001274	0. 94
	NT2RP3001297	F-NT2RP3001297	0.94
	NT2RP3001307	F-NT2RP3001307	0.87
	NT2RP3001325	F-NT2RP3001325	0.71
20	NT2RP3001338	F-NT2RP3001338	0.90
30	NT2RP3001355	F-NT2RP3001355	0.40
	NT2RP3001356	F-NT2RP3001356	0.88
	NT2RP3001384	F-NT2RP3001384	0. 91
	NT2RP3001392	F-NT2RP3001392	0.42
35	NT2RP3001396	F-NT2RP3001396	0. 56
	NT2RP3001398	F-NT2RP3001398	0. 94
	NT2RP3001399	F-NT2RP3001399	0. 94
	NT2RP3001407	F-NT2RP3001407	0. 54
40	NT2RP3001420	F-NT2RP3001420	0.74
	NT2RP3001426	F-NT2RP3001426	0. 92
	NT2RP3001427	F-NT2RP3001427	0.40
	NT2RP3001428	F-NT2RP3001428	0. 94
45	NT2RP3001447	F-NT2RP3001447	0. 57
	NT2RP3001449	F-NT2RP3001449	0. 62
	NT2RP3001453	F-NT2RP3001453	0.94
	NT2RP3001457	F-NT2RP3001457	0.89
50	NT2RP3001472	F-NT2RP3001472	0.82
	NT2RP3001490	F-NT2RP3001490	0. 53
	NT2RP3001495	F-NT2RP3001495	0.94
	NT2RP3001497	F-NT2RP3001497	0. 40
55	NT2RP3001529	F-NT2RP3001529	0.94
	NT2RP3001538	F-NT2RP3001538	0. 93

	NT-2RP3001554	F-NT2RP3001554	0. 91
	NT2RP3001587	F-NT2RP3001587	0.94
	NT2RP3001608	F-NT2RP3001608	0.94
5	NT2RP3001621	F-NT2RP3001621	
	NT2RP3001629	F-NT2RP3001629	0.42
	NT2RP3001642	F-NT2RP3001642	0.94
	NT2RP3001646	F-NT2RP3001646	0.94
10	NT2RP3001671	F-NT2RP3001671	0.77
	NT2RP3001672	F-NT2RP3001672	0.88
	NT2RP3001676	F-NT2RP3001676	0.88
	NT2RP3001678	F-NT2RP3001678	0. 44
15	NT2RP3001679	F-NT2RP3001679	0. 44
	NT2RP3001688	F-NT2RP3001688	0. 67
	NT2RP3001690	F-NT2RP3001690,	
	NT2RP3001698	F-NT2RP3001698	0.86
20	NT2RP3001708	F-NT2RP3001708	0. 38
20	NT2RP3001712	F-NT2RP3001712	0. 75
	NT2RP3001716	F-NT2RP3001716	0.77
	NT2RP3001724	F-NT2RP3001724	0. 65
0.5	NT2RP3001727	F-NT2RP3001727	0.84
25	NT2RP3001730	F-NT2RP3001730	0.77
	NT2RP3001739	F-NT2RP3001739	0. 78
	NT2RP3001753	F-NT2RP3001753	0.79
	NT2RP3001764	F-NT2RP3001764	0.33
30	NT2RP3001777	F-NT2RP3001777	0.87
	NT2RP3001792	F-NT2RP3001792	0.86
	NT2RP3001799	F-NT2RP3001799	0.79
	NT2RP3001819	F-NT2RP3001819	0.31
35	NT2RP3001844	F-NT2RP3001844	0.34
	NT2RP3001854	F-NT2RP3001854	0.94
	NT2RP3001855	F-NT2RP3001855	0.88
	NT2RP3001857	F-NT2RP3001857	0.41
40	NT2RP3001896	F-NT2RP3001896	0.94
	NT2RP3001915	F-NT2RP3001915	0.42
	NT2RP3001929	F-NT2RP3001929	0.49
	NT2RP3001931	F-NT2RP3001931	0.94
45	NT2RP3001938	F-NT2RP3001938	0.32
	NT2RP3001943	F-NT2RP3001943	0. 70
	NT2RP3001944	F-NT2RP3001944	0. 94
	NT2RP3001969	F-NT2RP3001969	0. 54
50	NT2RP3002007	F-NT2RP3002007	0. 56
	NT2RP3002014	F-NT2RP3002014	0. 94
	NT2RP3002033	F-NT2RP3002033	0. 94
	NT2RP3002045	F-NT2RP3002045	0. 91
55	NT2RP3002054	F-NT2RP3002054	0. 94
	NT2RP3002062	F-NT2RP3002062	0. 62

	NT2RP3002063	F-NT2RP3002063	0.34
	NT2RP3002081	F-NT2RP3002081	0. 57
	NT2RP3002097	F-NT2RP3002097	0.72
5	NT2RP3002102	F-NT2RP3002102	0. 90
	NT2RP3002108	F-NT2RP3002108	0.74
	NT2RP3002142	F-NT2RP3002142	0. 61
	NT2RP3002146	F-NT2RP3002146	0. 91
10	NT2RP3002147	F-NT2RP3002147	0. 37
	NT2RP3002151	F-NT2RP3002151	0. 90
	NT2RP3002163	F-NT2RP3002163	0.76
	NT2RP3002165	F-NT2RP3002165	0. 89
15	NT2RP3002181	F-NT2RP3002181	0. 78
	NT2RP3002244	F-NT2RP3002244	0.81
	NT2RP3002248	F-NT2RP3002248,	0.36
	NT2RP3002255	F-NT2RP3002255	0. 53
20	NT2RP3002273	F-NT2RP3002273	0.90
	NT2RP3002276	F-NT2RP3002276	0.39
	NT2RP3002303	F-NT2RP3002303	0.40
	NT2RP3002304	F-NT2RP3002304	0.44
25	NT2RP3002330	F-NT2RP3002330	0.94
20	NT2RP3002343	F-NT2RP3002343	0.94
	NT2RP3002351	F-NT2RP3002351	0.94
	NT2RP3002377	F-NT2RP3002377	0.52
30	NT2RP3002399	F-NT2RP3002399	0. 91
30	NT2RP3002402	F-NT2RP3002402	0.33
	NT2RP3002455	F-NT2RP3002455	0.42
	NT2RP3002484	F-NT2RP3002484	0.52
	NT2RP3002501	F-NT2RP3002501	0. 57
35	NT2RP3002512	F-NT2RP3002512	0. 63
	NT2RP3002529	F-NT2RP3002529	0. 74
	NT2RP3002545	F-NT2RP3002545	0. 94
	NT2RP3002549	F-NT2RP3002549	0.94
40	NT2RP3002566	F-NT2RP3002566	0.74
	NT2RP3002587	F-NT2RP3002587	0.94
	NT2RP3002590	F-NT2RP3002590	0. 37
	NT2RP3002602	F-NT2RP3002602	0.39
45	NT2RP3002603	F-NT2RP3002603	0.46
	NT2RP3002628	F-NT2RP3002628	0.31
	NT2RP3002631	F-NT2RP3002631	0.79
	NT2RP3002650 NT2RP3002659	F-NT2RP3002650	0.94
50		F-NT2RP3002659	0.93
	NT2RP3002660 NT2RP3002663	F-NT2RP3002660 F-NT2RP3002663	0. 46 0. 79
	NT2RP3002663		
	NT2RP3002671	F-NT2RP3002671 F-NT2RP3002682	0. 34 0. 52
55	NT2RP3002682	F-NT2RP3002682	0. 52
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	NT2RP3002701	F-NT2RP3002701	0.62
	NT2RP3002763	F-NT2RP3002763	0. 83
	NT2RP3002785	F-NT2RP3002785	0. 33
5	NT2RP3002818	F-NT2RP3002818	0. 91
	NT2RP3002861	F-NT2RP3002861	0.31
	NT2RP3002869	F-NT2RP3002869	0.47
	NT2RP3002876	F-NT2RP3002876	0. 58
10	NT2RP3002877	F-NT2RP3002877	0.41
	NT2RP3002909	F-NT2RP3002909	0. 52
	NT2RP3002911	F-NT2RP3002911	0.73
	NT2RP3002948	F-NT2RP3002948	0.60
15	NT2RP3002953	F-NT2RP3002953	0. 37
	NT2RP3002969	F-NT2RP3002969	0.47
	NT2RP3002972	F-NT2RP3002972,	0.87
	NT2RP3002985	F-NT2RP3002985	0. 55
20	NT2RP3002988	F-NT2RP3002988	0. 82
	NT2RP3003008	F-NT2RP3003008	0. 94
	NT2RP3003061	F-NT2RP3003061	0. 93
	NT2RP3003068	F-NT2RP3003068	0. 45
25	NT2RP3003071	F-NT2RP3003071	0.76
20	NT2RP3003078	F-NT2RP3003078	0. 40
	NT2RP3003101	F-NT2RP3003101	0. 41
	NT2RP3003139	F-NT2RP3003139	0.43
22	NT2RP3003145	F-NT2RP3003145	0. 94
30	NT2RP3003150	F-NT2RP3003150	0.94
	NT2RP3003157	F-NT2RP3003157	0. 81
	NT2RP3003185	F-NT2RP3003185	0. 87
	NT2RP3003193	F-NT2RP3003193	0. 94
35	NT2RP3003197	F-NT2RP3003197	0. 74
	NT2RP3003203	F-NT2RP3003203	0. 94
	NT2RP3003204	F-NT2RP3003204	0. 74
	NT2RP3003210	F-NT2RP3003210	0. 45
40	NT2RP3003212	F-NT2RP3003212	0. 51
	NT2RP3003230	F-NT2RP3003230	0. 94
	NT2RP3003242	F-NT2RP3003242	0. 92
	NT2RP3003251	F-NT2RP3003251	0. 93
45	NT2RP3003278	F-NT2RP3003278	0. 83
	NT2RP3003282	F-NT2RP3003282	0. 67
	NT2RP3003290	F-NT2RP3003290	0. 62
	NT2RP3003301	F-NT2RP3003301	0. 49
50	NT2RP3003302	F-NT2RP3003302	0. 32
	NT2RP3003311	F-NT2RP3003311	0. 34
	NT2RP3003313	F-NT2RP3003313	0. 88
	NT2RP3003327	F-NT2RP3003327	0.88
55	NT2RP3003330 NT2RP3003344	F-NT2RP3003330	0. 94
	N12NF30U3344	F-NT2RP3003344	0. 67

	NT2RP3003353	F-NT2RP3003353	0. 65
	NT2RP3003377	F-NT2RP3003377	0.87
	NT2RP3003385	F-NT2RP3003385	0.94
5	NT2RP3003409	F-NT2RP3003409	0.41
	NT2RP3003411	F-NT2RP3003411	0. 47
	NT2RP3003427	F-NT2RP3003427	0.33
	NT2RP3003490	F-NT2RP3003490	0. 56
10	NT2RP3003491	F-NT2RP3003491	0.37
	NT2RP3003543	F-NT2RP3003543	0.92
	NT2RP3003552	F-NT2RP3003552	0.64
	NT2RP3003555	F-NT2RP3003555	0.94
15	NT2RP3003564	F-NT2RP3003564	0.86
	NT2RP3003589	F-NT2RP3003589	0.88
	NT2RP3003621	F-NT2RP3003621,	0.49
	NT2RP3003656	F-NT2RP3003656	0. 77
20	NT2RP3003659	F-NT2RP3003659	0. 32
	NT2RP3003672	F-NT2RP3003672	0.49
	NT2RP3003680	F-NT2RP3003680	0. 94
	NT2RP3003686	F-NT2RP3003686	0. 78
25	NT2RP3003701	F-NT2RP3003701	0.36
	NT2RP3003716	F-NT2RP3003716	0. 34
	NT2RP3003726	F-NT2RP3003726	0. 72
	NT2RP3003795	F-NT2RP3003795	0.34
30	NT2RP3003805	F-NT2RP3003805	0. 74
	NT2RP3003809	F-NT2RP3003809	0. 73
	NT2RP3003825	F-NT2RP3003825	0. 94
	NT2RP3003831	F-NT2RP3003831	0. 52
35	NT2RP3003833	F-NT2RP3003833	0. 31
33	NT2RP3003846	F-NT2RP3003846	0. 54
	NT2RP3003870	F-NT2RP3003870	0. 91
	NT2RP3003876 NT2RP3003914	F-NT2RP3003876 F-NT2RP3003914	0. 42 0. 94
40	NT2RP3003914	F-NT2RP3003918	0. 94
40	NT2RP3003918	F-NT2RP3003932	0. 64
	NT2RP3003932	F-NT2RP3003992	0. 94
	NT2RP3004013	F-NT2RP3004013	0. 60
	NT2RP3004041	F-NT2RP3004041	0. 52
45	NT2RP3004051	F-NT2RP3004051	0. 75
	NT2RP3004078	F-NT2RP3004078	0. 73
	NT2RP3004093	F-NT2RP3004093	0. 64
	NT2RP3004125	F-NT2RP3004125	0. 94
50	NT2RP3004155	F-NT2RP3004155	0. 89
	NT2RP3004189	F-NT2RP3004189	0. 46
	NT2RP3004206	F-NT2RP3004206	0. 59
	NT2RP3004207	F-NT2RP3004207	0. 39
55	NT2RP3004209	F-NT2RP3004209	0. 54

	NT2RP3004242	F-NT2RP3004242	0.61
	NT2RP3004246	F-NT2RP3004246	0, 91
	NT2RP3004253	F-NT2RP3004253	0. 41
5	NT2RP3004258	F-NT2RP3004258	
	NT2RP3004262	F-NT2RP3004262	0.56
	NT2RP3004282	F-NT2RP3004282	0.90
	NT2RP3004332	F-NT2RP3004332	0. 53
10	NT2RP3004341	F-NT2RP3004341	0.40
	NT2RP3004348	F-NT2RP3004348	0. 94
	NT2RP3004378	F-NT2RP3004378	0.40
	NT2RP3004424	F-NT2RP3004424	0. 75
15	NT2RP3004428	F-NT2RP3004428	0. 85
	NT2RP3004451	F-NT2RP3004451	0.85
	NT2RP3004454	F-NT2RP3004454,	
	NT2RP3004466	F-NT2RP3004466	0.80
20	NT2RP3004472	F-NT2RP3004472	0.46
20	NT2RP3004480		0. 77
	NT2RP3004490	F-NT2RP3004490	0. 65
	NT2RP3004498	F-NT2RP3004498	0. 91
	NT2RP3004504	F-NT2RP3004504	0. 88.
25	NT2RP3004507	F-NT2RP3004507	0. 86
	NT2RP3004534	F-NT2RP3004534	0. 32
	NT2RP3004539	F-NT2RP3004539	0.76
	NT2RP3004544	F-NT2RP3004544	0.94
30	NT2RP3004566	F-NT2RP3004566	0.94
	NT2RP3004569	F-NT2RP3004569	0.69
	NT2RP3004572	F-NT2RP3004572	0. 65
	NT2RP3004578	F-NT2RP3004578	0. 52
35	NT2RP3004594	F-NT2RP3004594	0.39
	NT2RP3004617	F-NT2RP3004617	0.94
	NT2RP3004618	F-NT2RP3004618	0.41
	NT2RP3004669	F-NT2RP3004669	0.82
40	NT2RP3004670	F-NT2RP3004670	0.32
	NT2RP4000008	F-NT2RP4000008	0.94
	NT2RP4000051	F-NT2RP4000051	0.94
	NT2RP4000109	F-NT2RP4000109	0.84
45	NT2RP4000111	F-NT2RP4000111	0.49
	NT2RP4000129	F-NT2RP4000129	0.94
	NT2RP4000147	F-NT2RP4000147	0. 92
	NT2RP4000150	F-NT2RP4000150	0.41
50	NT2RP4000151	F-NT2RP4000151	0. 33
	NT2RP4000159	F-NT2RP4000159	0. 75
	NT2RP4000185	F-NT2RP4000185	0. 94
	NT2RP4000210	F-NT2RP4000210	0. 83
55	NT2RP4000212	F-NT2RP4000212	0. 94
	NT2RP4000243	F-NT2RP4000243	0. 94

	NT2RP4000246	F-NT2RP4000246	0. 59
	NT2RP4000259	F-NT2RP4000259	0.94
	NT2RP4000263	F-NT2RP4000263	0.62
5	NT2RP4000290	F-NT2RP4000290	0. 92
	NT2RP4000312	F-NT2RP4000312	0. 39
	NT2RP4000323	F-NT2RP4000323	0. 73
	NT2RP4000355	F-NT2RP4000355	0.94
10	NT2RP4000360	F-NT2RP4000360	0. 55
	NT2RP4000367	F-NT2RP4000367	0.60
	NT2RP4000370	F-NT2RP4000370	0.93
	NT2RP4000376	F-NT2RP4000376	0.88
15	NT2RP4000381	F-NT2RP4000381	0.39
	NT2RP4000398	F-NT2RP4000398	0.64
	NT2RP4000415	F-NT2RP4000415,	0.94
	NT2RP4000417	F-NT2RP4000417	0. 33
20	NT2RP4000448	F-NT2RP4000448	0. 33
	NT2RP4000449	F-NT2RP4000449	033
	NT2RP4000455	F-NT2RP4000455	0. 37
	NT2RP4000457	F-NT2RP4000457	0.36
25	NT2RP4000480	F-NT2RP4000480	0.88
•	NT2RP4000481	F-NT2RP4000481	0. 51
	NT2RP4000498	F-NT2RP4000498	0. 64
	NT2RP4000500	F-NT2RP4000500	0. 67
30	NT2RP4000518	F-NT2RP4000518	0. 34
	NT2RP4000524	F-NT2RP4000524	0. 94
	NT2RP4000528	F-NT2RP4000528	0. 65
	NT2RP4000541	F-NT2RP4000541	0. 44
35	NT2RP4000556	F-NT2RP4000556	0. 69
	NT2RP4000560	F-NT2RP4000560	0. 67
	NT2RP4000588 NT2RP4000614	F-NT2RP4000588 F-NT2RP4000614	0. 94 0. 94
	NT2RP4000614	F-NT2RP4000638	0. 74
40	NT2RP4000638	F-NT2RP4000638	0. 74
•	NT2RP4000657	F-NT2RP4000657	0. 94
	NT2RP4000704	F-NT2RP4000704	0. 69
	NT2RP4000713	F-NT2RP4000713	0. 93
45	NT2RP4000724	F-NT2RP4000724	0, 32
45	NT2RP4000728	F-NT2RP4000728	0. 34
	NT2RP4000737	F-NT2RP4000737	0. 48
	NT2RP4000739	F-NT2RP4000739	0. 40
50	NT2RP4000781	F-NT2RP4000781	0. 94
50	NT2RP4000787	F-NT2RP4000787	0.94
	NT2RP4000817	F-NT2RP4000817	0. 60
	NT2RP4000833	F-NT2RP4000833	0. 35
	NT2RP4000837	F-NT2RP4000837	0.80
55	NT2RP4000839	F-NT2RP4000839	0.33

	NT2RP4000855	F-NT2RP4000855	0. 74
	NT2RP4000865	F-NT2RP4000865	0.77
	NT2RP4000878	F-NT2RP4000878	0. 79
5	NT2RP4000879	F-NT2RP4000879	0. 41
	NT2RP4000907	F-NT2RP4000907	0. 44
	NT2RP4000918	F-NT2RP4000918	0. 88
	NT2RP4000925	F-NT2RP4000925	0. 89
10	NT2RP4000927	F-NT2RP4000927	0.81
	NT2RP4000928	F-NT2RP4000928	0. 39
	NT2RP4000929	F-NT2RP4000929	0. 34
	NT2RP4000955	F-NT2RP4000955	0. 77
15	NT2RP4000973	F-NT2RP4000973	0.36
	NT2RP4000975	F-NT2RP4000975	0. 87
	NT2RP4000979	F-NT2RP4000979,	0.50
	NT2RP4000984	F-NT2RP4000984	0. 66
20	NT2RP4000989	F-NT2RP4000989	0. 94
	NT2RP4000997	F-NT2RP4000997	0. 68
	NT2RP4001004	F-NT2RP4001004	0.40
	NT2RP4001006	F-NT2RP4001006	0. 91
25	NT2RP4001010	F-NT2RP4001010	0.64
	NT2RP4001029	F-NT2RP4001029	0.77
	NT2RP4001041	F-NT2RP4001041	0. 79
	NT2RP4001057	F-NT2RP4001057	0.65
30	NT2RP4001064	F-NT2RP4001064	0. 94
30	NT2RP4001078	F-NT2RP4001078	0. 39
	NT2RP4001079	F-NT2RP4001079	0. 94
	NT2RP4001080	F-NT2RP4001080	0. 58
25	NT2RP4001086	F-NT2RP4001086	0. 78
35	NT2RP4001095	F-NT2RP4001095	0. 80
	NT2RP4001100	F-NT2RP4001100	0. 79
	NT2RP4001117		0. 60
	NT2RP4001122		0. 82
40	NT2RP4001126	F-NT2RP4001126	0. 55
	NT2RP4001138	F-NT2RP4001138	0.94
	NT2RP4001143		0. 41
	NT2RP4001148	F-NT2RP4001148	0.37
45	NT2RP4001149	F-NT2RP4001149	0.50
	NT2RP4001150 NT2RP4001159	F-NT2RP4001150 F-NT2RP4001159	0. 94 0. 78
	NT2RP4001174	F-NT2RP4001174	0.78
	NT2RP4001174	F-NT2RP4001174	0. 80
50	NT2RP4001200	F-NT2RP4001200	0.38
	NT2RP4001207	F-NT2RP4001210	0. 38
	NT2RP4001213	F-NT2RP4001210	0. 46
	NT2RP4001213	F-NT2RP4001219	0. 38
55	NT2RP4001219	F-NT2RP4001219	0. 61
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	NT2RP4001235	F-NT2RP4001235	0.88
	NT2RP4001256	F-NT2RP4001256	0.60
_	NT2RP4001260	F-NT2RP4001260	0. 94
5	NT2RP4001274	F-NT2RP4001274_	0.82
	NT2RP4001276	F-NT2RP4001276	0.68
	NT2RP4001313	F-NT2RP4001313	0.77
	NT2RP4001315	F-NT2RP4001315	0. 87
10	NT2RP4001336	F-NT2RP4001336	0. 48
	NT2RP4001339	F-NT2RP4001339	0. 41
	NT2RP4001343	F-NT2RP4001343	0.63
	NT2RP4001345	F-NT2RP4001345	0.89
15	NT2RP4001351	F-NT2RP4001351	0. 87
	NT2RP4001353	F-NT2RP4001353	0.72
	NT2RP4001372	F-NT2RP4001372,	0. 78
	NT2RP4001373	F-NT2RP4001373	0.60
20	NT2RP4001375	F-NT2RP4001375	0.43
	NT2RP4001379	F-NT2RP4001379	0.77
	NT2RP4001389	F-NT2RP4001389	0.41
	NT2RP4001414	F-NT2RP4001414	0.90
25	NT2RP4001433	F-NT2RP4001433	0. 39
	NT2RP4001442	F-NT2RP4001442	0.59
	NT2RP4001447	F-NT2RP4001447	0. 33
	NT2RP4001474	F-NT2RP4001474	0.89
30	NT2RP4001483	F-NT2RP4001483	0.94
	NT2RP4001498	F-NT2RP4001498	0. 59
	NT2RP4001502	F-NT2RP4001502	0. 61
	NT2RP4001507	F-NT2RP4001507	0.91
35	NT2RP4001524	F-NT2RP4001524	0. 61
	NT2RP4001529	F-NT2RP4001529	0.75
	NT2RP4001547	F-NT2RP4001547	0.77
	NT2RP4001551	F-NT2RP4001551	0.79
40	NT2RP4001555	F-NT2RP4001555	0.79
40	NT2RP4001567 NT2RP4001568	F-NT2RP4001567	0.39
	NT2RP4001508	F-NT2RP4001568	0. 45
	NT2RP4001571	F-NT2RP4001571	0. 76
_	NT2RP4001575	F-NT2RP4001574 F-NT2RP4001575	0.91
45	NT2RP4001575		0.57
	NT2RP4001610	F-NT2RP4001592 F-NT2RP4001610	0.71
	NT2RP4001614	F-NT2RP4001614	0. 94
	NT2RP4001634	F-NT2RP4001634	0. 73 0. 40
50	NT2RP4001638	F-NT2RP4001638	0. 37
	NT2RP4001644	F-NT2RP4001644	0. 94
	NT2RP4001656	F-NT2RP4001656	0. 92
	NT2RP4001677	F-NT2RP4001677	0. 47
55	NT2RP4001679	F-NT2RP4001679	0. 54
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	NT2RP4001696	F-NT2RP4001696	0. 62
	NT2RP4001725	F-NT2RP4001725	0. 94
_	NT2RP4001730	F-NT2RP4001730	0.94
5	NT2RP4001739	F-NT2RP4001739_	0.82
	NT2RP4001753	F-NT2RP4001753	0.66
	NT2RP4001760	F-NT2RP4001760	0.39
	NT2RP4001790	F-NT2RP4001790	0.64
10	NT2RP4001803	F-NT2RP4001803	0.87
	NT2RP4001822	F-NT2RP4001822	0.84
	NT2RP4001823	F-NT2RP4001823	0.56
	NT2RP4001828	F-NT2RP4001828	0. 45
15	NT2RP4001838	F-NT2RP4001838	0. 87
	NT2RP4001841	F-NT2RP4001841	0. 58
	NT2RP4001849	F-NT2RP4001849	0. 88
	NT2RP4001861	F-NT2RP4001861	0.80
20	NT2RP4001893	F-NT2RP4001893	0.94
	NT2RP4001896	F-NT2RP4001896	0. 93
	NT2RP4001901	F-NT2RP4001901	0.89
	NT2RP4001927	F-NT2RP4001927	0. 77
25	NT2RP4001938	F-NT2RP4001938	0. 94
	NT2RP4001946	F-NT2RP4001946	
	NT2RP4001950	F-NT2RP4001950	0. 45
	NT2RP4001953	F-NT2RP4001953	0. 49
30	NT2RP4001966	F-NT2RP4001966	0.86
	NT2RP4001975	F-NT2RP4001975	0.94
	NT2RP4002018	F-NT2RP4002018	0. 67
	NT2RP4002047	F-NT2RP4002047	0. 67
35	NT2RP4002052 NT2RP4002058	F-NT2RP4002052	0. 42
	NT2RP4002058	F-NT2RP4002058 F-NT2RP4002071	0. 78
	NT2RP4002071	F-NT2RP4002071	0. 94 0. 38
	NT2RP4002078	F-NT2RP4002078	0. 55
40	NT2RP4002083	F-NT2RP4002083	0.86
	NT2RP4002408	F-NT2RP4002408	0.64
	NT2RP4002791	F-NT2RP4002791	0.84
	NT2RP4002888	F-NT2RP4002888	0.31
45	NT2RP5003459	F-NT2RP5003459	0. 94
	NT2RP5003461	F-NT2RP5003461	0. 94
	NT2RP5003477	F-NT2RP5003477	0. 44
	NT2RP5003492	F-NT2RP5003492	0. 57
50	NT2RP5003500	F-NT2RP5003500	0.94
	NT2RP5003506	F-NT2RP5003506	0.94
	NT2RP5003512	F-NT2RP5003512	0. 42
	NT2RP5003522	F-NT2RP5003522	0.94
55	NT2RP5003524	F-NT2RP5003524	0. 58
55	NT2RP5003534	F-NT2RP5003534	0. 32

	OVARCIUUUUUI	F-0VARC1000001	0. 74
	OVARC1000004	F-0VARC1000004	0. 90
	0VARC1000006	F-0VARC1000006	0. 94
5	OVARC1000013	F-0VARC1000013	0. 48
	OVARC1000014	F-0VARC1000014	0. 69
	OVARC1000035	F-0VARC1000035	0. 59
	OVARC1000060	F-0VARC1000060	0. 76
10	0VARC1000087	F-0VARC1000087	0. 31
	0VARC1000091	F-0VARC1000091	0. 88
	OVARC1000106	F-0VARC1000106	0. 88
	OVARC1000109	F-0VARC1000109	0. 94
15	OVARC1000113	F-0VARC1000113	0. 40
	0VARC1000114	F-0VARC1000114	0. 53
	OVARC1000139	F-0VARC1000139,	0. 88
	0VARC1000148	F-0VARC1000148	0. 94
20	0VARC1000151		0. 34
	0VARC1000168	F-0VARC1000168	0. 56
	0VARC1000209	F-0VARC1000209	0. 94
	0VARC1000212	F-0VARC1000212	0. 87
25	0VARC1000241	F-0VARC1000241	0. 59
20	0VARC1000288	F-0VARC1000288	0. 87
	0VARC1000304	F-0VARC1000304	0. 94
	OVARC1000309	F-0VARC1000309	0. 76
20	0VARC1000321	F-0VARC1000321	0.90
30	0VARC1000326	F-0VARC1000326	0. 46
	0VARC1000335	F-0VARC1000335	0. 55
	0VARC1000347	F-0VARC1000347	0. 43
	0VARC1000384	F-0VARC1000384	0. 90
35	0VARC1000408	F-0VARC1000408	0. 42
	0VARC1000411	F-0VARC1000411	0. 65
	0VARC1000420		0. 84
	0VARC1000437		0. 94
40			0. 94
		F-0VARC1000442	0. 94
	0VARC1000443	F-0VARC1000443	0. 74
	0VARC1000461	F-0VARC1000461	0. 89
45	OVARC1000465	F-0VARC1000465	0. 75
	OVARC1000466	F-0VARC1000466	0. 94
	0VARC1000473	F-0VARC1000473	0. 31
	0VARC1000479	F-0VARC1000479	0. 42
50	0VARC1000520	F-0VARC1000520	0. 94
	0VARC1000556	F-0VARC1000556	0. 94
	0VARC1000557	F-0VARC1000557	0. 72
	0VARC1000564	F-0VARC1000564	0. 94
55	0VARC1000576	F-0VARC1000576	0. 40
	OVARC1000588	F-0VARC1000588	0. 45

	OVARC1000605	F-0VARC1000605	0.58
	0VARC1000622	F-0VARC1000622	0.80
	OVARC1000640	F-0VARC1000640	0. 35
5	OVARC1000649	F-0VARC1000649	0.94
	OVARC1000661	F-0VARC1000661	0.87
	0VARC1000681	F-0VARC1000681	0.84
	OVARC1000682	F-0VARC1000682	0.38
10	0VARC1000689	F-0VARC1000689	0.36
	0VARC1000703	F-0VARC1000703	0. 92
	0VARC1000722	F-0VARC1000722	0.94
	0VARC1000730	F-0VARC1000730	0. 90
15	0VARC1000746	F-0VARC1000746	0.42
	0VARC1000771	F-0VARC1000771	0. 94
	0VARC1000781	F-0VARC1000781,	
	OVARC1000787	F-0VARC1000787	0. 41
20	OVARC1000800	F-0VARC1000800	0. 85
	0VARC1000834	F-0VARC1000834	0.62
	0VARC1000846	F-0VARC1000846	0.49
	0VARC1000850	F-0VARC1000850	0.94
25	0VARC1000862	F-0VARC1000862	0.54
	0VARC1000876	F-0VARC1000876	0.74
	0VARC1000883	F-0VARC1000883	0. 55
	0VARC1000885	F-0VARC1000885	0.94
20	0VARC1000886	F-0VARC1000886	0.61
30	OVARC1000890	F-0VARC1000890	0.45
	0VARC1000912	F-0VARC1000912	0.63
	OVARC1000915	F-0VARC1000915	0.31
	0VARC1000924	F-0VARC1000924	0.82
35	0VARC1000945	F-0VARC1000945	0.74
	0VARC1000959	F-0VARC1000959	0. 33
	0VARC1000964	F-0VARC1000964	0. 60
	0VARC1000984	F-0VARC1000984	0. 59
40	0VARC1000996	F-0VARC1000996	0. 83
	0VARC1000999	F-0VARC1000999	0. 94
	0VARC1001004	F-0VARC1001004	0. 69
	0VARC1001010	F-0VARC1001010	0. 51
45	0VARC1001032	F-0VARC1001032	0. 32
	0VARC1001034	F-0VARC1001034	0. 78
	0VARC1001038	F-0VARC1001038	0. 93
	0VARC1001040	F-0VARC1001040	0. 68
50	0VARC1001044	F-0VARC1001044	0. 66
	0VARC1001055	F-0VARC1001055	0. 76
	OVARC1001065	F-0VARC1001065	0. 31
	0VARC1001068	F-0VARC1001068	0. 63
55	0VARC1001074	F-0VARC1001074	0.82
	0VARC1001092	F-0VARC1001092	0. 50

	OVARC1001107	F-0VARC1001107	0.65
	OVARC1001113	F-0VARC1001113	0.86
	OVARC1001154	F-0VARC1001154	0. 43
5	OVARC1001161	F-0VARC1001161	0.75
	OVARC1001162	F-0VARC1001162	0.61
	OVARC1001167	F-0VARC1001167	0.31
	OVARC1001170	F-0VARC1001170	0. 46
10	OVARC1001171	F-0VARC1001171	0.78
	OVARC1001173	F-0VARC1001173	0.71
	OVARC1001176	F-0VARC1001176	0.83
	OVARC1001180	F-0VARC1001180	0.53
15	OVARC1001188	F-0VARC1001188	0. 42
	OVARC1001200	F-0VARC1001200	0. 94
	OVARC1001232	F-0VARC1001232,	
	OVARC1001243	F-0VARC1001243	0. 59
22	OVARC1001244	F-0VARC1001244	0. 79
20	OVARC1001270	F-0VARC1001270	0. 48
	OVARC1001271	F-0VARC1001271	0. 67
	OVARC1001296	F-0VARC1001296	0. 60
	OVARC1001306	F-0VARC1001306	0. 44
25	OVARC1001329	F-0VARC1001329	0. 91
	OVARC1001341	F-0VARC1001341	0.82
	OVARC1001344	F-0VARC1001344	0. 94
	OVARC1001360	F-0VARC1001360	0.94
30	OVARC1001369	F-0VARC1001369	0.36
	OVARC1001372	F-0VARC1001372	0.63
	OVARC1001376	F-0VARC1001376	0.81
	OVARC1001381	F-0VARC1001381	0. 78
35	OVARC1001391	F-0VARC1001391	0. 33
	OVARC1001399	F-0VARC1001399	0. 67
	OVARC1001417	F-0VARC1001417	0. 94
	OVARC1001419	F-0VARC1001419	0. 34
40	OVARC1001425	F-0VARC1001425	0. 73
	OVARC1001436	F-0VARC1001436	0. 59
	OVARC1001453	F-0VARC1001453	0. 45
	OVARC1001476	F-0VARC1001476	0. 85
45	OVARC1001489	F-0VARC1001489	0. 35
	OVARC1001496	F-0VARC1001496	0. 90
	OVARC1001506	F-0VARC1001506	0. 59
	OVARC1001525	F-0VARC1001525	0.87
50	OVARC1001555	F-0VARC1001555	0. 67
	0VARC1001577	F-0VARC1001577	0.94
	OVARC1001610	F-0VARC1001610	0.38
	OVARC1001611	F-0VARC1001611	0.34
55	OVARC1001702	F-0VARC1001702	0.88
	OVARC1001703	F-0VARC1001703	0. 76

	OVARC1001711	F-0VARC1001711	0.82
	OVARC1001713	F-0VARC1001713	0.82
	OVARC1001731	F-0VARC1001731	0.94
5	0VARC1001762	F-0VARC1001762	
	OVARC1001766	F-0VARC1001766	0.94
	OVARC1001767	F-0VARC1001767	0.70
	OVARC1001768	F-0VARC1001768	0. 32
10	OVARC1001791	F-0VARC1001791	0.74
	OVARC1001802	F-0VARC1001802	0.76
	OVARC1001809		0.63
	OVARC1001820		0. 03
_	OVARC1001828		0. 78
15	0VARC1001828		
			0.94
	OVARC1001873		
	OVARC1001879	•	0.94
20	0VARC1001880		0.46
	OVARC1001916		0.82
	0VARC1001942		0.72
	OVARC1001943		0.49
25	0VARC1001949	F-0VARC1001949	0.46
	OVARC1001950	F-0VARC1001950	0.94
		F-0VARC1001987	0. 91
	0VARC1002050	F-0VARC1002050	0. 85
30	0VARC1002082		0.94
30	OVARC1002107		
	0VARC1002112		0. 94
	OVARC1002127		0. 80
		F-0VARC1002138	0. 46
35	0VARC1002156		0. 43
	0VARC1002158		0. 94
	0VARC1002165		0. 88
	0VARC1002182		0. 94
40	PLACE1000004	F-PLACE1000004	0. 43
	PLACE1000005	F-PLACE1000005	0. 37
	PLACE1000007	F-PLACE1000007	0. 72
	PLACE1000014	F-PLACE1000014	0. 38
45	PLACE1000048	F-PLACE1000048	0. 36
	PLACE1000050	F-PLACE1000050	0. 94
	PLACE1000061	F-PLACE1000061	0. 56
	PLACE1000066	F-PLACE1000066	0. 94
50	PLACE1000081	F-PLACE1000081	0.37
50	PLACE1000133	F-PLACE1000133	0.53
	PLACE1000142	F-PLACE1000142	0.94
	PLACE1000184	F-PLACE1000184	0. 73
	PLACE1000185	F-PLACE1000185	0.86
55	PLACE1000213	F-PLACE1000213	0. 34

	PLACE1000236	F-PLACE1000236	0. 76
	PLACE1000246	F-PLACE1000246	0. 91
	PLACE1000308	F-PLACE1000308	0.40
5	PLACE1000347	F-PLACE1000347	
	PLACE1000374	F-PLACE1000374	0. 58
	PLACE1000380	F-PLACE1000380	0. 94
	PLACE1000383	F-PLACE1000383	0. 39
10	PLACE1000401	F-PLACE1000401	0. 94
	PLACE1000406	F-PLACE1000406	0. 94
	PLACE1000420	F-PLACE1000420	0. 82
	PLACE1000435	F-PLACE1000435	0. 32
15	PLACE1000444	F-PLACE1000444	0. 41
	PLACE1000453	F-PLACE1000453	0. 76
	PLACE1000492	F-PLACE1000492,	
	PLACE1000547	F-PLACE1000547	0. 94
00	PLACE1000562	F-PLACE1000562	0.94
20	PLACE1000564	F-PLACE1000564	0. 45
	PLACE1000588	F-PLACE1000588	0. 67
	PLACE1000596	F-PLACE1000596	0. 60
	PLACE1000610	F-PLACE1000610	0. 94
25	PLACE1000611	F-PLACE1000611	0. 48
	PLACE1000636	F-PLACE1000636	0. 92
	PLACE1000653	F-PLACE1000653	0. 57
	PLACE1000656	F-PLACE1000656	0. 94
30	PLACE1000706	F-PLACE1000706	0. 94
	PLACE1000712	F-PLACE1000712	0. 78
	PLACE1000716	F-PLACE1000716	0. 94
	PLACE1000748	F-PLACE1000748	0. 52
35	PLACE1000755	F-PLACE1000755	0.87
	PLACE1000769	F-PLACE1000769	0.94
	PLACE1000785	F-PLACE1000785	0. 33
	PLACE1000786	F-PLACE1000786	0.88
40	PLACE1000793	F-PLACE1000793	0. 79
	PLACE1000798	F-PLACE1000798	0.59
	PLACE1000849	F-PLACE1000849	0.64
	PLACE1000856	F-PLACE1000856	0.40
45	PLACE1000863	F-PLACE1000863	0. 94
	PLACE1000909	F-PLACE1000909	0.94
	PLACE1000948	F-PLACE1000948	0. 92
	PLACE1000972	F-PLACE1000972	0. 73
50	PLACE1000977	F-PLACE1000977	0. 72
	PLACE1000987	F-PLACE1000987	0. 93
	PLACE1001000	F-PLACE1001000	0. 37
	PLACE1001010	F-PLACE1001010	0. 50
22	PLACE1001024	F-PLACE1001024	0.94
55	PLACE1001036	F-PLACE1001036	0. 51

	PLACE1001054	F-PLACE1001054	0. 88
	PLACE1001062	F-PLACE1001062	0. 57
	PLACE1001092	F-PLACE1001092	0. 94
5	PLACE1001104	F-PLACE1001104_	
	PLACE1001168	F-PLACE1001168	0. 41
	PLACE1001171	F-PLACE1001171	0.36
	PLACE1001185	F-PLACE1001185	0.93
10	PLACE1001238	F-PLACE1001238	0.87
	PLACE1001241	F-PLACE1001241	0.67
	PLACE1001257	F-PLACE1001257	0.40
	PLACE1001280	F-PLACE1001280	0.70
15	PLACE1001294	F-PLACE1001294	0.49
	PLACE1001304	F-PLACE1001304	0.94
	PLACE1001311	F-PLACE1001311,	0.53
	PLACE1001351	F-PLACE1001351	0.86
20	PLACE1001366	F-PLACE1001366	0.39
20	PLACE1001383	F-PLACE1001383	0.94
	PLACE1001387	F-PLACE1001387	0.34
	PLACE1001399	F-PLACE1001399	0.38
	PLACE1001412	F-PLACE1001412	0.94
25	PLACE1001456	F-PLACE1001456	0.42
	PLACE1001484	F-PLACE1001484	0.88
	PLACE1001503	F-PLACE1001503	0.85
	PLACE1001545	F-PLACE1001545	0.92
30	PLACE1001551	F-PLACE1001551	0.39
	PLACE1001570	F-PLACE1001570	0.40
	PLACE1001602	F-PLACE1001602	0.60
	PLACE1001608	F-PLACE1001608	0.40
35	PLACE1001610	F-PLACE1001610	0.43
	PLACE1001632	F-PLACE1001632	0.48
	PLACE1001634	F-PLACE1001634	0.60
	PLACE1001692	F-PLACE1001692	0.74
40	PLACE1001729	F-PLACE1001729	0.69
	PLACE1001739	F-PLACE1001739	0.49
	PLACE1001740	F-PLACE1001740	0.50
	PLACE1001761	F-PLACE1001761	0.71
45	PLACE1001771	F-PLACE1001771	0.39
45	PLACE1001781	F-PLACE1001781	0.57
	PLACE1001810	F-PLACE1001810	0. 57
	PLACE1001817	F-PLACE1001817	0. 91
	PLACE1001844	F-PLACE1001844	0.34
50	PLACE1001869	F-PLACE1001869	0.34
	PLACE1001912	F-PLACE1001912	0. 82
	PLACE1001920	F-PLACE1001920	0. 85
•	PLACE1001928	F-PLACE1001928	0.60
55	PLACE1001983	F-PLACE1001983	0. 39

	PLACE1001989	F-PLACE1001989	0:74
	PLACE1002004	F-PLACE1002004	0.70
	PLACE1002046	F-PLACE1002046	0.81
5	PLACE1002072	F-PLACE1002072	
	PLACE1002073	F-PLACE1002073	0. 70
	PLACE1002090	F-PLACE1002090	0.64
	PLACE1002115	F-PLACE1002115	0.60
10	PLACE1002119	F-PLACE1002119	0. 71
	PLACE1002140	F-PLACE1002140	0.94
	PLACE1002143	F-PLACE1002163	0.56
	PLACE1002170	F-PLACE1002170	0.60
	PLACE1002170	F-PLACE1002171	0. 72
15			0. 72
	PLACE1002213	F-PLACE1002213	
	PLACE1002342	F-PLACE1002342,	
	PLACE1002395	F-PLACE1002395	0.87
20	PLACE1002433	F-PLACE1002433	0. 73
	PLACE1002437	F-PLACE1002437	0.46
	PLACE1002438	F-PLACE1002438	0.41
	PLACE1002450	F-PLACE1002450	0. 48
25	PLACE1002465	F-PLACE1002465	0.34
	PLACE1002474	F-PLACE1002474	0.89
	PLACE1002499	F-PLACE1002499	0. 94
	PLACE1002529	F-PLACE1002529	0. 45
30	PLACE1002532	F-PLACE1002532	0. 76
30	PLACE1002571	F-PLACE1002571	0.40
	PLACE1002625	F-PLACE1002625	0.36
	PLACE1002655	F-PLACE1002655	0.41
	PLACE1002665	F-PLACE1002665	0.80
35	PLACE1002685	F-PLACE1002685	0. 31
	PLACE1002714	F-PLACE1002714	0. 90
	PLACE1002722	F-PLACE1002722	0. 43
	PLACE1002775	F-PLACE1002775	0.94
40	PLACE1002794	F-PLACE1002794	0. 90
	PLACE1002811	F-PLACE1002811	0.88
	PLACE1002815	F-PLACE1002815	0.40
	PLACE1002816	F-PLACE1002816	0. 73
45	PLACE1002834	F-PLACE1002834	0. 41
	PLACE1002839	F-PLACE1002839	0.73
	PLACE1002851	F-PLACE1002851	0.90
	PLACE1002908	F-PLACE1002908	0.94
	PLACE1002941	F-PLACE1002941	0.33
50	PLACE1002991	F-PLACE1002991	0.39
	PLACE1002993	F-PLACE1002993	0. 94
	PLACE1002996	F-PLACE1002996	0.40
	PLACE1003025	F-PLACE1003025	0. 77
55	PLACE1003027	F-PLACE1003027	0.54

	PLACE1003044	F-PLACE1003044	0.71
	PLACE1003045	F-PLACE1003045	0.89
	PLACE1003092	F-PLACE1003092	0.59
5	PLACE1003100	F-PLACE1003100	0.38
	PLACE1003108	F-PLACE1003108	0.94
	PLACE1003136	F-PLACE1003136	0. 87
	PLACE1003145	F-PLACE1003145	0. 94
10	PLACE1003174	F-PLACE1003174	0. 46
	PLACE1003176	F-PLACE1003176	0. 48
	PLACE1003190	F-PLACE1003190	0. 57
	PLACE1003200	F-PLACE1003200	0. 47
45	PLACE1003249	F-PLACE1003249	0. 33
15	PLACE1003256	F-PLACE1003256	0. 57
	PLACE1003296	F-PLACE1003296,	
	PLACE1003302	F-PLACE1003302	0. 94
	PLACE1003334	F-PLACE1003334	0. 78
20	PLACE1003342	F-PLACE1003342	0. 74
	PLACE1003353	F-PLACE1003353	0.71
	PLACE1003366	F-PLACE1003366	0.66
	PLACE1003369	F-PLACE1003369	0. 91
25	PLACE1003383	F-PLACE1003383	0.70
	PLACE1003394	F-PLACE1003394	0.94
	PLACE1003454	F-PLACE1003454	0.94
	PLACE1003493	F-PLACE1003493	0.93
30	PLACE1003537	F-PLACE1003537	0.94
	PLACE1003553	F-PLACE1003553	0.34
	PLACE1003592	F-PLACE1003592	0. 33
	PLACE1003596	F-PLACE1003596	0.89
35	PLACE1003602	F-PLACE1003602	0. 94
	PLACE1003605	F-PLACE1003605	0.94
	PLACE1003611	F-PLACE1003611	0.80
	PLACE1003625	F-PLACE1003625	0.39
40	PLACE1003669	F-PLACE1003669	0. 79
	PLACE1003704	F-PLACE1003704	0. 43
	PLACE1003709	F-PLACE1003709	0. 90
	PLACE1003711	F-PLACE1003711	0. 74
45	PLACE1003723	F-PLACE1003723	0. 63
40	PLACE1003738	F-PLACE1003738	0. 91
	PLACE1003762	F-PLACE1003762	0. 94
	PLACE1003771	F-PLACE1003771	0. 56
	PLACE1003783	F-PLACE1003783	0.39
50	PLACE1003784	F-PLACE1003784	0.61
	PLACE1003795	F-PLACE1003795	0. 32
	PLACE1003870	F-PLACE1003870	0. 94
	PLACE1003885	F-PLACE1003885	0. 94
55	PLACE1003886	F-PLACE1003886	0. 94

	PLACE1003888	F-PLACE1003888	0.36
	PLACE1003892	F-PLACE1003892	0. 92
	PLACE1003903	F-PLACE1003903	0. 68
5	PLACE1003915	F-PLACE1003915	0. 54
	PLACE1003923	F-PLACE1003923	0. 52
	PLACE1003936	F-PLACE1003936	0. 67
	PLACE1003968	F-PLACE1003968	0. 59
10	PLACE1004103	F-PLACE1004103	0. 94
	PLACE1004104	F-PLACE1004104	0. 44
	PLACE1004114	F-PLACE1004114	0. 91
	PLACE1004114	F-PLACE1004128	0. 72
15	PLACE1004149	F-PLACE1004149	0. 72
	PLACE1004149	F-PLACE1004156	0. 50
	PLACE1004150	F-PLACE1004161,	
	PLACE1004181	•	
	PLACE1004183	F-PLACE1004183	0.62
20		F-PLACE1004197	0.39
	PLACE1004203	F-PLACE1004203	0.94
	PLACE1004258	F-PLACE1004258	0.46
	PLACE1004270	F-PLACE1004270	0.65
25	PLACE1004277	F-PLACE1004277	0.86
	PLACE1004289	F-PLACE1004289	0. 79
	PLACE1004302	F-PLACE1004302	0.57
	PLACE1004316	F-PLACE1004316	0. 64
30	PLACE1004358	F-PLACE1004358	0. 85
	PLACE1004376	F-PLACE1004376	0. 64
	PLACE1004388	F-PLACE1004388	0. 94
	PLACE1004405	F-PLACE1004405	0. 82
	PLACE1004428	F-PLACE1004428	0. 68
35	PLACE1004437	F-PLACE1004437	0. 94
	PLACE1004451	F-PLACE1004451	0. 39
	PLACE1004460	F-PLACE1004460	0. 92
	PLACE1004473	F-PLACE1004473	0. 94
40	PLACE1004510	F-PLACE1004510	0. 94
	PLACE1004516	F-PLACE1004516	0. 58
	PLACE1004548	F-PLACE1004548	0. 59
	PLACE1004550	F-PLACE1004550	0. 94
45	PLACE1004564	F-PLACE1004564	0. 53
	PLACE1004629	F-PLACE1004629	0. 34
		F-PLACE1004645	0. 59
	PLACE1004646	F-PLACE1004646	0. 35
50	PLACE1004664	F-PLACE1004664	0.31
SU	PLACE1004672	F-PLACE1004672	0.83
	PLACE1004674	F-PLACE1004674	0.90
	PLACE1004686	F-PLACE1004686	0. 31
	PLACE1004691	F-PLACE1004691	0.89
55	PLACE1004722	F-PLACE1004722	0. 48

	PLACE1004736	F-PLACE1004736	0. 80
	PLACE1004740	F-PLACE1004740	0. 34
	PLACE1004743	F-PLACE1004743	0. 70
5	PLACE1004751	F-PLACE1004751	0. 49
	PLACE1004777	F-PLACE1004777	0. 72
	PLACE1004804	F-PLACE1004804	0.46
	PLACE1004814	F-PLACE1004814	0. 70
10	PLACE1004824	F-PLACE1004824	0. 63
	PLACE1004868	F-PLACE1004868	0. 94
	PLACE1004885	F-PLACE1004885	0. 63
	PLACE1004902	F-PLACE1004902	0, 56
15	PLACE1004918	F-PLACE1004918	0. 85
	PLACE1004930	F-PLACE1004930	0, 83
	PLACE1004937	F-PLACE1004937,	0. 46
	PLACE1004969	F-PLACE1004969	0. 62
20	PLACE1004982	F-PLACE1004982	0.61
	PLACE1005026	F-PLACE1005026	0.81
	PLACE1005027	F-PLACE1005027	0.91
	PLACE1005046	F-PLACE1005046	0. 31
25 .	PLACE1005055	F-PLACE1005055	0. 57
	PLACE1005066	F-PLACE1005066	0.68
	PLACE1005101	F-PLACE1005101	0.94
	PLACE1005102	F-PLACE1005102	0.94
•	PLACE1005181	F-PLACE1005181	0.94
30	PLACE1005187	F-PLACE1005187	0. 94
	PLACE1005206	F-PLACE1005206	0.67
	PLACE1005232	F-PLACE1005232	0.72
	PLACE1005243	F-PLACE1005243	0.81
35	PLACE1005261	F-PLACE1005261	0.75
	PLACE1005266	F-PLACE1005266	0. 55
	PLACE1005277	F-PLACE1005277	0.43
	PLACE1005287	F-PLACE1005287	0.77
40	PLACE1005305	F-PLACE1005305	0.94
	PLACE1005308	F-PLACE1005308	0.46
	PLACE1005313	F-PLACE1005313	0. 94
	PLACE1005327	F-PLACE1005327	0. 82
45	PLACE1005331	F-PLACE1005331	0. 94
	PLACE1005335	F-PLACE1005335	0. 94
	PLACE1005373	F-PLACE1005373	0.71
	PLACE1005374	F-PLACE1005374	0. 82
50	PLACE1005480	F-PLACE1005480	0. 43
	PLACE1005481	F-PLACE1005481	0. 42
	PLACE1005494	F-PLACE1005494	0. 34
	PLACE1005530	F-PLACE1005530	0. 94
55	PLACE1005550	F-PLACE1005550	0. 86
55	PLACE1005554	F-PLACE1005554	0. 34

	PLACE1005557	F-PLACE1005557	0.80
	PLACE1005595	F-PLACE1005595	0.69
•	PLACE1005603	F-PLACE1005603	0. 92
5	PLACE1005623	F-PLACE1005623	0. 68
	PLACE1005630	F-PLACE1005630	0.58
	PLACE1005646	F-PLACE1005646	0.31
	PLACE1005656	F-PLACE1005656	0.85
10	PLACE1005698	F-PLACE1005698	0.91
	PLACE1005727	F-PLACE1005727	0.94
	PLACE1005739	F-PLACE1005739	0.87
	PLACE1005755	F-PLACE1005755	0.32
15	PLACE1005763	F-PLACE1005763	0.74
	PLACE1005799	F-PLACE1005799	0. 47
	PLACE1005803	F-PLACE1005803,	0.88
	PLACE1005804	F-PLACE1005804	0. 36
20	PLACE1005813	F-PLACE1005813	0. 76
	PLACE1005828	F-PLACE1005828	0. 43
	PLACE1005834	F-PLACE1005834	0.89
	PLACE1005851	F-PLACE1005851	0. 39
25	PLACE1005876	F-PLACE1005876	0.51
	PLACE1005890	F-PLACE1005890	0. 81
	PLACE1005921	F-PLACE1005921	0. 90
	PLACE1005923	F-PLACE1005923	0. 70
30	PLACE1005925	F-PLACE1005925	0. 55
	PLACE1005934	F-PLACE1005934	0. 94
	PLACE1005936	F-PLACE1005936	0. 41
	PLACE1005951	F-PLACE1005951	0.34
35	PLACE1005953	F-PLACE1005953	0.31
	PLACE1005955 PLACE1005966	F-PLACE1005955 F-PLACE1005966	0. 89 0. 73
	PLACE1005980 PLACE1005990	F-PLACE1005900	0.73
	PLACE1005990	F-PLACE1005990	0. 94
40	PLACE1006037	F-PLACE1006037	0.65
•	PLACE1006040	F-PLACE1006040	0. 94
	PLACE1006119	F-PLACE1006119	
	PLACE1006139	F-PLACE1006139	0.94
	PLACE1006157	F-PLACE1006157	0.34
45	PLACE1006159	F-PLACE1006159	0. 50
	PLACE1006167	F-PLACE1006167	0. 94
	PLACE1006170	F-PLACE1006170	0. 94
	PLACE1006195	F-PLACE1006195	0. 31
50	PLACE1006196	F-PLACE1006196	0. 84
	PLACE1006225	F-PLACE1006225	0.87
	PLACE1006236	F-PLACE1006236	0. 92
	PLACE1006239	F-PLACE1006239	0. 53
55	PLACE1006246	F-PLACE1006246	0. 44

	PLACE1006248	F-PLACE1006248	0.49
	PLACE1006325	F-PLACE1006325	0.74
	PLACE1006335	F-PLACE1006335	0.82
5	PLACE1006357	F-PLACE1006357_	0.33
	PLACE1006385	F-PLACE1006385	0. 54
	PLACE1006412	F-PLACE1006412	0. 79
	PLACE1006414	F-PLACE1006414	0. 52
10	PLACE1006438	F-PLACE1006438	0. 94
	PLACE1006445	F-PLACE1006445	0.37
	PLACE1006469	F-PLACE1006469	0. 85
	PLACE1006482	F-PLACE1006482	0. 94
15	PLACE1006488	F-PLACE1006488	0.94
	PLACE1006492	F-PLACE1006492	0. 32
	PLACE1006531	F-PLACE1006531;	0. 45
	PLACE1006552	F-PLACE1006552	0. 42
20	PLACE1006598	F-PLACE1006598	0. 33
	PLACE1006615	F-PLACE1006615	0. 89
	PLACE1006626	F-PLACE1006626	0. 69
·	PLACE1006673	F-PLACE1006673	0.61
25	PLACE1006678	F-PLACE1006678	0. 53
	PLACE1006704	F-PLACE1006704	0. 47
	PLACE1006731	F-PLACE1006731	0. 72
	PLACE1006782	F-PLACE1006782	0. 68
30	PLACE1006819	F-PLACE1006819	0. 57
	PLACE1006829	F-PLACE1006829	0. 47
	PLACE1006883 PLACE1006901	F-PLACE1006883	0.88
	PLACE1006901	F-PLACE1006901 F-PLACE1006917	0. 43 0. 56
35	PLACE1006917	F-PLACE1006917	0. 45
•	PLACE1006935	F-PLACE1006935	0. 43
	PLACE1006956	F-PLACE1006956	0. 77
	PLACE1006958	F-PLACE1006958	0. 75
40	PLACE1006961	F-PLACE1006961	0. 34
40	PLACE1006962	F-PLACE1006962	0. 94
	PLACE1006966	F-PLACE1006966	
	PLACE1006989	F-PLACE1006989	0. 47
-	PLACE1007014	F-PLACE1007014	0.94
45	PLACE1007021	F-PLACE1007021	0.74
	PLACE1007053	F-PLACE1007053	0.73
	PLACE1007068	F-PLACE1007068	0.43
	PLACE1007105	F-PLACE1007105	0.88
50	PLACE1007112	F-PLACE1007112	0.82
	PLACE1007178	F-PLACE1007178	0. 91
	PLACE1007226	F-PLACE1007226	0. 79
	PLACE1007238	F-PLACE1007238	0.94
55	PLACE1007239	F-PLACE1007239	0.94

•	PLACE1007242	F-PLACE1007242	0. 48
	PLACE1007243	F-PLACE1007243	0.77
	PLACE1007257	F-PLACE1007257	0. 54
5	PLACE1007274	F-PLACE1007274	0. 94
	PLACE1007282	F-PLACE1007282	0.94
	PLACE1007301	F-PLACE1007301	0.34
	PLACE1007317	F-PLACE1007317	0.43
10	PLACE1007342	F-PLACE1007342	0.72
	PLACE1007346	F-PLACE1007346	0.92
	PLACE1007367	F-PLACE1007367	0.43
	PLACE1007375	F-PLACE1007375	0.85
15	PLACE1007386	F-PLACE1007386	0.58
	PLACE1007402	F-PLACE1007402	0.74
	PLACE1007409	F-PLACE1007409,	0.39
	PLACE1007416	F-PLACE1007416	0.80
20	PLACE1007450	F-PLACE1007450	0.41
	PLACE1007452	F-PLACE1007452	0.68
	PLACE1007454	F-PLACE1007454	0.39
	PLACE1007460	F-PLACE1007460	0.58
25	PLACE1007484	F-PLACE1007484	0.94
	PLACE1007488	F-PLACE1007488	0.50
	PLACE1007507	F-PLACE1007507	0.44
	PLACE1007511	F-PLACE1007511	0.86
30	PLACE1007524	F-PLACE1007524	0.66
30	PLACE1007537	F-PLACE1007537	0.93
	PLACE1007544	F-PLACE1007544	0.94
	PLACE1007547	F-PLACE1007547	0.65
	PLACE1007583	F-PLACE1007583	0.87
35	PLACE1007598	F-PLACE1007598	0.39
	PLACE1007618	F-PLACE1007618	0.62
	PLACE1007621	F-PLACE1007621	0.63
	PLACE1007632 PLACE1007645	F-PLACE1007632 F-PLACE1007645	0.75 0.94
40	PLACE1007649	F-PLACE1007649	0. 56
	PLACE1007649	F-PLACE1007688	0. 94
	PLACE1007690	F-PLACE1007690	0. 44
	PLACE1007697	F-PLACE1007697	0. 93
45	PLACE1007705	F-PLACE1007705	0.61
	PLACE1007706	F-PLACE1007706	0.47
	PLACE1007725	F-PLACE1007725	0.61
	PLACE1007729	F-PLACE1007729	0. 53
50	PLACE1007730	F-PLACE1007730	0.31
	PLACE1007746	F-PLACE1007746	0. 81
	PLACE1007791	F-PLACE1007791	0. 46
	PLACE1007810	F-PLACE1007810	0. 73
55	PLACE1007843	F-PLACE1007843	0. 31

	PLACE1007846	F-PLACE1007846	0.38
	PLACE1007858	F-PLACE1007858	0.79
	PLACE1007897	F-PLACE1007897	0.94
5	PLACE1007908	F-PLACE1007908	0.94
	PLACE1007946	F-PLACE1007946	0. 52
	PLACE1007954	F-PLACE1007954	0.43
	PLACE1007955	F-PLACE1007955	0.41
10	PLACE1007958	F-PLACE1007958	0.37
	PLACE1007969	F-PLACE1007969	0. 87
	PLACE1007990	F-PLACE1007990	0.94
	PLACE1008000	F-PLACE1008000	0.94
15	PLACE1008002	F-PLACE1008002	0.93
	PLACE1008044	F-PLACE1008044	0.72
	PLACE1008095	F-PLACE1008095,	0.94
	PLACE1008122	F-PLACE1008122	0.94
20	PLACE1008129	F-PLACE1008129	0.80
	PLACE1008132	F-PLACE1008132	0. 37
	PLACE1008177	F-PLACE1008177	0. 74
	PLACE1008198	F-PLACE1008198	0.94
25	PLACE1008209	F-PLACE1008209	0.54
	PLACE1008273	F-PLACE1008273	0.94
	PLACE1008275	F-PLACE1008275	0.82
	PLACE1008280	F-PLACE1008280	0.34
30	PLACE1008309	F-PLACE1008309	0.90
	PLACE1008329 PLACE1008356	F-PLACE1008329 F-PLACE1008356	0. 38 0. 93
	PLACE1008368	F-PLACE1008368	0. 43
	PLACE1008398	F-PLACE1008398	0.78
35	PLACE1008401	F-PLACE1008401	0.70
	PLACE1008402	F-PLACE1008402	0.88
	PLACE1008429	F-PLACE1008429	0.39
	PLACE1008457	F-PLACE1008457	0.77
40	PLACE1008465	F-PLACE1008465	0.94
	PLACE1008488	F-PLACE1008488	0.42
	PLACE1008524	F-PLACE1008524	0.41
	PLACE1008531	F-PLACE1008531	0.94
45	PLACE1008532	F-PLACE1008532	0.61
	PLACE1008533	F-PLACE1008533	0. 45
	PLACE1008568	F-PLACE1008568	0.65
	PLACE1008603	F-PLACE1008603	0.42
50	PLACE1008626	F-PLACE1008626	0. 94
50	PLACE1008627	F-PLACE1008627	0.72
	PLACE1008629	F-PLACE1008629	0. 33
	PLACE1008650	F-PLACE1008650	0. 79
	PLACE1008693	F-PLACE1008693	0. 32
55	PLACE1008696	F-PLACE1008696	0. 39

	PLACE1008790	F-PLACE1008790	0. 81
	PLACE1008808	F-PLACE1008808	0.51
	PLACE1008813	F-PLACE1008813	0.36
5	PLACE1008854	F-PLACE1008854	
	PLACE1008867	F-PLACE1008854	0.34
	PLACE1008887	F-PLACE1008887	0. 49
	PLACE1008807	F-PLACE1008902	0. 49
10	PLACE1008902 PLACE1008925		
		F-PLACE1008925 F-PLACE1008934	0.94
	PLACE1008934 PLACE1009027		0.70
		F-PLACE1009027	0. 32
4-	PLACE1009045	F-PLACE1009045	0.42
15	PLACE1009060	F-PLACE1009060	0.69
	PLACE1009090	F-PLACE1009090	0.46
	PLACE1009091	F-PLACE1009091,	
	PLACE1009094	F-PLACE1009094	0. 44
20	PLACE1009099	F-PLACE1009099	0. 56
	PLACE1009110	F-PLACE1009110	0.84
	PLACE1009111	F-PLACE1009111	0.80
	PLACE1009113	F-PLACE1009113	0.88
25	PLACE1009130	F-PLACE1009130	0.60
	PLACE1009158	F-PLACE1009158	0.41
	PLACE1009166	F-PLACE1009166	0.94
	PLACE1009174 PLACE1009186	F-PLACE1009174	0.38
30	PLACE1009186	F-PLACE1009186 F-PLACE1009190	0.59
	PLACE1009190 PLACE1009230	F-PLACE1009190	0. 66 0. 87
	PLACE1009230	F-PLACE1009230	0. 67
	PLACE1009319	F-PLACE1009319	0. 53
35	PLACE1009328	F-PLACE1009328	0. 66
	PLACE1009335	F-PLACE1009335	0. 47
	PLACE1009338	F-PLACE1009338	0. 70
	PLACE1009368	F-PLACE1009368	0.94
40	PLACE1009375	F-PLACE1009375	0.76
40	PLACE1009388	F-PLACE1009388	0. 46
	PLACE1009404	F-PLACE1009404	0.72
	PLACE1009434	F-PLACE1009434	0.50
	PLACE1009443	F-PLACE1009443	0. 90
45	PLACE1009444	F-PLACE1009444	0. 33
	PLACE1009459	F-PLACE1009459	0. 94
	PLACE1009468	F-PLACE1009468	0. 88
	PLACE1009476	F-PLACE1009476	0. 84
50	PLACE1009524	F-PLACE1009524	0.88
	PLACE1009542	F-PLACE1009542	0. 55
	PLACE1009571	F-PLACE1009571	0. 61
	PLACE1009581	F-PLACE1009581	0. 94
55	PLACE1009596	F-PLACE1009596	0.44
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	PLACE1009607	F-PLACE1009607	0. 47
	PLACE1009622	F-PLACE1009622	0.65
	PLACE1009659	F-PLACE1009659	0.36
5	PLACE1009665	F-PLACE1009665	0. 43
	PLACE1009670	F-PLACE1009670	0. 33
	PLACE1009708	F-PLACE1009708	0.80
	PLACE1009721	F-PLACE1009721	0.81
10	PLACE1009731	F-PLACE1009731	0.47
	PLACE1009763	F-PLACE1009763	0.94
	PLACE1009794	F-PLACE1009794	0.33
	PLACE1009798	F-PLACE1009798	0.34
15	PLACE1009845	F-PLACE1009845	0.66
	PLACE1009861	F-PLACE1009861	0. 72
	PLACE1009886	F-PLACE1009886,	
	PLACE1009908	F-PLACE1009908	0. 92
•	PLACE1009921	F-PLACE1009921	0. 70
-	PLACE1009971	F-PLACE1009971	0.40
	PLACE1009992	F-PLACE1009992	0.89
	PLACE1009995	F-PLACE1009995	0.88
	PLACE1009997	F-PLACE1009997	0. 41
25	PLACE1010023	F-PLACE1010023	0.90
	PLACE1010031	F-PLACE1010031	0. 71
	PLACE1010053	F-PLACE1010053	0. 51
	PLACE1010074	F-PLACE1010074	0. 75
30	PLACE1010076	F-PLACE1010076	0.53
	PLACE1010096	F-PLACE1010096	0. 44
	PLACE1010102	F-PLACE1010102	0.77
	PLACE1010105	F-PLACE1010105	0. 58
35	PLACE1010106	F-PLACE1010106	0. 39
	PLACE1010134	F-PLACE1010134	0.85
	PLACE1010148	F-PLACE1010148	0. 32
	PLACE1010152	F-PLACE1010152	0.94
40	PLACE1010194	F-PLACE1010194	0.94
	PLACE1010202	F-PLACE1010202	0.33
	PLACE1010261	F-PLACE1010261	0.36
	PLACE1010274	F-PLACE1010274	0.94
45	PLACE1010293	F-PLACE1010293	0.55
	PLACE1010310	F-PLACE1010310	0. 41
	PLACE1010321	F-PLACE1010321	0.40
	PLACE1010324	F-PLACE1010324	0.40
50	PLACE1010329	F-PLACE1010329	0.36
50	PLACE1010362	F-PLACE1010362	0.34
	PLACE1010364	F-PLACE1010364	0. 37
	PLACE1010383	F-PLACE1010383	0.94
	PLACE1010401	F-PLACE1010401	0.53
55	PLACE1010481	F-PLACE1010481	0. 74

	PLACE1010491	F-PLACE1010491	0. 31
	PLACE1010492	F-PLACE1010492	0. 39
	PLACE1010522	F-PLACE1010522	0. 92
5	PLACE1010529	F-PLACE1010529	0. 65
	PLACE1010547	F-PLACE1010547	0. 82
	PLACE1010580	F-PLACE1010580	0. 72
	PLACE1010599	F-PLACE1010599	0. 83
10	PLACE1010616	F-PLACE1010616	0. 54
	PLACE1010622	F-PLACE1010622	0. 92
	PLACE1010629	F-PLACE1010629	0. 94
	PLACE1010630	F-PLACE1010630	0. 37
15	PLACE1010661	F-PLACE1010661	0. 56
	PLACE1010714	F-PLACE1010714	0. 68
	PLACE1010720	F-PLACE1010720,	0. 90
	PLACE1010743	F-PLACE1010743	0. 75
20	PLACE1010761	F-PLACE1010761	0. 69
20	PLACE1010771	F-PLACE1010771	0.94
	PLACE1010786	F-PLACE1010786	0. 80
	PLACE1010800	F-PLACE1010800	0.41
	PLACE1010811	F-PLACE1010811	0. 94
25	PLACE1010833	F-PLACE1010833	0.94
	PLACE1010856	F-PLACE1010856	0. 37
	PLACE1010857	F-PLACE1010857	0. 54
	PLACE1010870	F-PLACE1010870	0.50
30	PLACE1010877	F-PLACE1010877	0. 77
	PLACE1010900	F-PLACE1010900	0. 70
	PLACE1010917	F-PLACE1010917	0.94
	PLACE1010925	F-PLACE1010925	0. 79
35	PLACE1010926	F-PLACE1010926	0. 94
	PLACE1010942	F-PLACE1010942	0. 36
	PLACE1010944	F-PLACE1010944	0.61
	PLACE1010954	F-PLACE1010954	0. 83
40	PLACE1010960	F-PLACE1010960	0.38
	PLACE1010965	F-PLACE1010965	0.68
	PLACE1011026	F-PLACE1011026	0.38
	PLACE1011041	F-PLACE1011041	0.94
45	PLACE1011046	F-PLACE1011046	0.85
	PLACE1011054	F-PLACE1011054	0. 65
	PLACE1011056	F-PLACE1011056	0.82
	PLACE1011090	F-PLACE1011090	0.94
50	PLACE1011114 PLACE1011160	F-PLACE1011114 F-PLACE1011160	0.94
	PLACE1011160	F-PLACE10111100	0. 62 0. 40
	PLACE1011214 PLACE1011219	F-PLACE1011219	0. 48
	PLACE1011219	F-PLACE1011219	0. 48
ee	PLACE1011221	F-PLACE1011229	0. 46
55	FEMOL 1011229	F-FLAGE 1011229	U. 40

	PLACE1011263	F-PLACE1011263	0. 94
	PLACE1011273	F-PLACE1011273	0.32
	PLACE1011291	F-PLACE1011291	0. 55
5	PLACE1011310	F-PLACE1011310	0.74
	PLACE1011325	F-PLACE1011325	0. 39
	PLACE1011332	F-PLACE1011332	0.74
	PLACE1011371	F-PLACE1011371	0. 32
10	PLACE1011399	F-PLACE1011399	0.89
	PLACE1011433	F-PLACE1011433	0. 33
	PLACE1011472	F-PLACE1011472	0. 59
	PLACE1011477	F-PLACE1011477	0.80
15	PLACE1011563	F-PLACE1011563	0.83
	PLACE1011586	F-PLACE1011586	0. 82
	PLACE1011635	F-PLACE1011635,	0. 62
	PLACE1011646	F-PLACE1011646	0. 37
20	PLACE1011664	F-PLACE1011664	0. 65
20	PLACE1011675	F-PLACE1011675	0. 89
	PLACE1011682	F-PLACE1011682	0.81
	PLACE1011725	F-PLACE1011725	0.86
	PLACE1011783	F-PLACE1011783	0. 78
25	PLACE1011858	F-PLACE1011858	0.76
	PLACE1011875	F-PLACE1011875	0.49
	PLACE1011891	F-PLACE1011891	0.34
	PLACE1011896	F-PLACE1011896	0.34
30	PLACE1011922	F-PLACE1011922	0.46
	PLACE1011923	F-PLACE1011923	0.74
	PLACE1011982	F-PLACE1011982	0.71
	PLACE1012031	F-PLACE1012031	0.80
35	PLACE2000006	F-PLACE2000006	0.61
	PLACE2000007	F-PLACE2000007	0. 37
	PLACE2000014	F-PLACE2000014	0. 67
	PLACE2000015	F-PLACE2000015	0. 53
40	PLACE2000021	F-PLACE2000021	0.36
	PLACE2000030	F-PLACE2000030	0. 83
	PLACE2000033	F-PLACE2000033	0. 44
	PLACE2000034	F-PLACE2000034	0. 88
45	PLACE2000039	F-PLACE2000039	0.67
	PLACE2000047	F-PLACE2000047	0.84
	PLACE2000050	F-PLACE2000050	0. 41
	PLACE2000062	F-PLACE2000062	0.94
50	PLACE2000072	F-PLACE2000072	0.86
	PLACE2000097	F-PLACE2000097	0. 94
	PLACE2000100	F-PLACE2000100	0. 75
	PLACE2000111	F-PLACE2000111	0. 62
	PLACE2000124	F-PLACE2000124	0. 77
55	PLACE2000164	F-PLACE2000164	0. 57

	PLACE2000170	F-PLACE2000170	0.60
	PLACE2000172	F-PLACE2000172	0. 55
	PLACE2000223	F-PLACE2000223	0.54
5	PLACE2000235	F-PLACE2000235	0. 94
	PLACE2000246	F-PLACE2000246	0. 94
	PLACE2000274	F-PLACE2000274	0.82
	PLACE2000302	F-PLACE2000302	0. 92
10	PLACE2000341	F-PLACE2000341	0. 58
	PLACE2000359	F-PLACE2000359	0. 33
	PLACE2000371	F-PLACE2000371	0. 71
	PLACE2000373	F-PLACE2000373	0. 65
15	PLACE2000379	F-PLACE2000399	0. 41
19	PLACE2000393	F-PLACE2000404	0. 94
	PLACE2000404	F-PLACE2000404	
	PLACE2000411	F-PLACE2000417	0. 60
20	PLACE2000433	F-PLACE2000433	0.61
	PLACE2000438	F-PLACE2000438	0.56
	PLACE2000458	F-PLACE2000458	0. 82
	PLACE3000009	F-PLACE3000009	0.56
25	PLACE3000020	F-PLACE3000020	0.94
	PLACE3000029	F-PLACE3000029	0. 94
	PLACE3000059	F-PLACE3000059	0. 51
	PLACE3000070	F-PLACE3000070	0. 67
30	PLACE3000103	F-PLACE3000103	0. 34
30	PLACE3000121	F-PLACE3000121	0. 32
	PLACE3000142	F-PLACE3000142	0. 67
	PLACE3000145	F-PLACE3000145	0. 84
	PLACE3000147	F-PLACE3000147	0. 57
35	PLACE3000148	F-PLACE3000148	0. 94
	PLACE3000155	F-PLACE3000155	0. 42
	PLACE3000156	F-PLACE3000156	0. 40
	PLACE3000157	F-PLACE3000157	0. 75
40	PLACE3000160	F-PLACE3000160	0. 44
	PLACE3000194	F-PLACE3000194	0. 37
	PLACE3000197	F-PLACE3000197	0. 37
	PLACE3000221	F-PLACE3000221	0. 49
45	PLACE3000226	F-PLACE3000226	0. 39
	PLACE3000242	F-PLACE3000242	0. 94
	PLACE3000244	F-PLACE3000244	0. 56
	PLACE3000254	F-PLACE3000254	0. 60
	PLACE3000304	F-PLACE3000304	0. 42
50	PLACE3000310	F-PLACE3000310	0. 71
	PLACE3000320	F-PLACE3000320	0. 37
	PLACE3000322	F-PLACE3000322	0. 46
	PLACE3000331	F-PLACE3000331	0. 31
55	PLACE3000339	F-PLACE3000339	0. 45

	PLACE3000350	F-PLACE3000350	0. 52
	PLACE3000352	F-PLACE3000352	0. 65
	PLACE3000353	F-PLACE3000353	0.71
5	PLACE3000363	F-PLACE3000363	
	PLACE3000405	F-PLACE3000405	0. 52
	PLACE3000413	F-PLACE3000413	0. 85
	PLACE3000416	F-PLACE3000416	0.94
10	PLACE3000425	F-PLACE3000425	0.40
	PLACE3000455	F-PLACE3000455	0. 45
	PLACE3000477	F-PLACE3000477	0. 79
	PLACE4000009	F-PLACE4000009	0. 74
15	PLACE4000014	F-PLACE4000014	0. 46
	PLACE4000034	F-PLACE4000034	0. 89
	PLACE4000049	F-PLACE4000049,	0. 68
	PLACE4000052	F-PLACE4000052	0.80
20	PLACE4000063	F-PLACE4000063	0.69
	PLACE4000089	F-PLACE4000089	0. 44
	PLACE4000106	F-PLACE4000106	0. 34
	PLACE4000128	F-PLACE4000128	0.89
25	PLACE4000129	F-PLACE4000129	0. 61
	PLACE4000131	F-PLACE4000131	0. 55
	PLACE4000147	F-PLACE4000147	0.94
	PLACE4000156	F-PLACE4000156	0.37
30	PLACE4000192	F-PLACE4000192	0.88
30	PLACE4000211	F-PLACE4000211	0. 64
•	PLACE4000222	F-PLACE4000222	0.40
	PLACE4000230	F-PLACE4000230	0. 49
	PLACE4000259	F-PLACE4000259	0.84
35	PLACE4000261	F-PLACE4000261	0. 42
	PLACE4000269	F-PLACE4000269	0.60
	PLACE4000270	F-PLACE4000270	0.61
	PLACE4000300	F-PLACE4000300	0.85
40	PLACE4000323 PLACE4000326	F-PLACE4000323	0.76
	PLACE4000326	F-PLACE4000326 F-PLACE4000369	0. 86 0. 87
	PLACE4000389	F-PLACE4000387	0. 87
	PLACE4000387	F-PLACE4000387	0. 39
45	PLACE4000332	F-PLACE4000332	0. 88
	PLACE4000445	F-PLACE4000445	0. 40
	PLACE4000450	F-PLACE4000450	0. 89
	PLACE4000465	F-PLACE4000465	0. 38
50	PLACE4000487	F-PLACE4000487	0.44
	PLACE4000489	F-PLACE4000489	0. 51
	PLACE4000521	F-PLACE4000521	0.38
	PLACE4000522	F-PLACE4000522	0. 33
55	PLACE4000558	F-PLACE4000558	0.44
			• •

	PLACE4000581	F-PLACE4000581	0.59
	PLACE4000590	F-PLACE4000590	0.38
	PLACE4000593	F-PLACE4000593	0.33
5	PLACE4000612	F-PLACE4000612	0.37
	PLACE4000638	F-PLACE4000638	0. 38
	PLACE4000650	F-PLACE4000650	0. 59
	PLACE4000654	F-PLACE4000654	0.57
10	PLACE4000670	F-PLACE4000670	0.71
	SKNMC1000011	F-SKNMC1000011	0.94
	SKNMC1000013	F-SKNMC1000013	0.66
	SKNMC1000046	F-SKNMC1000046	0.72
15	SKNMC1000050	F-SKNMC1000050	0.47
	SKNMC1000091	F-SKNMC1000091	0.68
	THYR01000017	F-THYR01000017,	0.93
	THYR01000034	F-THYR01000034	0. 48
20	THYR01000040	F-THYR01000040	0. 32
	THYR01000070	F-THYR01000070	0. 75
	THYR01000072	F-THYR01000072	0.94
	THYR01000107	F-THYR01000107	0.94
25	THYR01000121	F-THYR01000121	0.46
25	THYR01000124	F-THYR01000124	0.37
	THYR01000163	F-THYR01000163	0. 45
	THYR01000173	F-THYR01000173	0.94
	THYR01000197	F-THYR01000197	0.78
30	THYR01000199	F-THYR01000199	0.55
	THYR01000206	F-THYR01000206	0. 59
	THYR01000242	F-THYR01000242	0.69
	THYR01000253	F-THYR01000253	0. 48
35	THYR01000270	F-THYR01000270	0.85
	THYRO1000288	F-THYR01000288	0.31
	THYR01000320	F-THYR01000320	0.94
	THYR01000327	F-THYR01000327	0.44
40	THYR01000343	F-THYR01000343	0. 82
	THYR01000358	F-THYR01000358	0.89
	THYR01000368	F-THYR01000368	0.94
	THYR01000381	F-THYR01000381	0.64
45	THYR01000387	F-THYR01000387	0.57
	THYR01000394	F-THYR01000394	0.94
	THYR01000395	F-THYR01000395	0.91
	THYR01000401	F-THYR01000401	0.73
50	THYR01000488	F-THYR01000488	0.94
	THYR01000501	F-THYR01000501	0.74
	THYR01000502	F-THYR01000502	0.64
	THYR01000505	F-THYR01000505	0.47
55	THYR01000569	F-THYR01000569	0.76
00	THYR01000570	F-THYR01000570	0. 79

	THYR01000585	F-THYR01000585	0. 60
	THYR01000605	F-THYR01000605	0.79
	THYR01000637	F-THYR01000637	0.80
5	THYR01000662	F-THYR01000662	
	THYR01000666	F-THYR01000666	0.47
	THYR01000676	F-THYR01000676	0. 94
	THYR01000684	F-THYR01000684	0. 72
10	THYR01000712	F-THYR01000712	0.94
	THYR01000715	F-THYR01000715	0. 94
	THYR01000734	F-THYR01000734	0. 44
	THYR01000756	F-THYR01000756	0. 94
15	THYR01000777	F-THYR01000777	0. 73
	THYR01000783	F-THYR01000783	0. 74
	THYR01000787	F-THYR01000787,	
	THYR01000829	F-THYR01000829	0. 43
••	THYR01000855	F-THYR01000855	0. 91
20	THYR01000916	F-THYR01000916	0. 64
	THYR01000926	F-THYR01000926	0. 39
	THYR01000934	F-THYR01000934	0. 94
	THYR01000951	F-THYR01000951	0. 94
25	THYR01000952	F-THYR01000952	0. 67
	THYR01000983	F-THYR01000983	0. 87
	THYR01000984	F-THYR01000984	0. 94
	THYR01000988	F-THYR01000988	0. 37
30	THYR01001033	F-THYR01001033	0. 74
	THYR01001100	F-THYR01001100	0.74
	THYR01001120	F-THYR01001120	0. 87
	THYR01001134	F-THYR01001134	0, 93
35	THYR01001142	F-THYR01001142	0.61
	THYR01001173	F-THYR01001173	0.50
	THYR01001189	F-THYR01001189	0.40
	THYR01001204	F-THYR01001204	0.94
40	THYR01001213	F-THYR01001213	0. 94
	THYR01001271	F-THYR01001271	0. 93
	THYR01001287	F-THYR01001287	0. 32
·	THYR01001313	F-THYR01001313	0. 53
45	THYR01001320	F-THYR01001320	0.68
	THYR01001347	F-THYR01001347	0.68
	THYR01001363	F-THYR01001363	0. 39
	THYR01001374	F-THYR01001374	0.94
50	THYR01001403	F-THYR01001403	0.49
	THYR01001405	F-THYR01001405	0. 90
	THYR01001406	F-THYR01001406	0.67
	THYR01001458	F-THYR01001458	0. 94
	THYR01001487	F-THYR01001487	0. 63
55	THYR01001537	F-THYR01001537	0. 94

	TIND01001504	E TINDO4004504	
	THYR01001584	F-THYR01001584	0.77
	THYR01001602	F-THYR01001602	0.64
_	THYR01001605	F-THYR01001605	0. 52
5	THYR01001637	F-THYR01001637_	.0,.68
	THYR01001656	F-THYR01001656	0. 82
	THYR01001661	F-THYR01001661	0.94
	THYR01001671	F-THYR01001671	0.94
10	THYR01001703	F-THYR01001703	0.87
	THYR01001721	F-THYR01001721	0.38
	THYR01001738	F-THYR01001738	0.51
	THYR01001793	F-THYR01001793	0.84
15	THYR01001809	F-THYR01001809	0. 73
•	THYR01001828	F-THYR01001828	0.88
	THYR01001854	F-THYR01001854,	0.47
	VESEN1000122	F-VESEN1000122	0.36
	Y79AA1000013	F-Y79AA1000013	0.94
20	Y79AA1000037	F-Y79AA1000037	0.94
	Y79AA1000059	F-Y79AA1000059	0.94
	Y79AA1000131	F-Y79AA1000131	0.56
	Y79AA1000181	F-Y79AA1000181	0.59
25	Y79AA1000202	F-Y79AA1000202	0.45
	Y79AA1000214	F-Y79AA1000214	0.93
	Y79AA1000230	F-Y79AA1000230	0. 94
	Y79AA1000231	F-Y79AA1000231	0.36
30	Y79AA1000258	F-Y79AA1000258	0. 36
	Y79AA1000268	F-Y79AA1000268	0.92
	Y79AA1000313	F-Y79AA1000313	0.91
	Y79AA1000328	F-Y79AA1000328	0.94
35	Y79AA1000342	F-Y79AA1000342	0.83
33	Y79AA1000349	F-Y79AA1000349	0.50
	Y79AA1000355	F-Y79AA1000355	0. 94
	Y79AA1000368	F-Y79AA1000368	0. 91
	Y79AA1000420	F-Y79AA1000420	0. 94
40	Y79AA1000469	F-Y79AA1000469	0.51
	Y79AA1000480	F-Y79AA1000480	0. 94
	Y79AA1000539	F-Y79AA1000539	0.74
	Y79AA1000540	F-Y79AA1000540	0.94
45	Y79AA1000560	F-Y79AA1000560	0.93
	Y79AA1000574	F-Y79AA1000574	0. 87
	Y79AA1000589	F-Y79AA1000589	0. 94
	Y79AA1000627	F-Y79AA1000627	0. 55
50	Y79AA1000705	F-Y79AA1000705	0.76
	Y79AA1000734	F-Y79AA1000734	0.60
	Y79AA1000748	F-Y79AA1000748	0. 45
	Y79AA1000752	F-Y79AA1000752	0. 94
55	Y79AA1000774	F-Y79AA1000774	0.38
55			

	Y79AA1000782	F-Y79AA1000782	0. 47
	Y79AA1000784	F-Y79AA1000784	0.93
5	Y79AA1000794	F-Y79AA1000794	0.67
•	Y79AA1000800	F-Y79AA10008QQ_	_0.,94
	Y79AA1000824	F-Y79AA1000824	0.90
	Y79AA1000827	F-Y79AA1000827	0.94
	Y79AA1000833	F-Y79AA1000833	0.34
10	Y79AA1000850	F-Y79AA1000850	0. 92
	Y79AA1000962	F-Y79AA1000962	0.69
	Y79AA1000966	F-Y79AA1000966	0.64
	Y79AA1000968	F-Y79AA1000968	0.64
15	Y79AA1000976	F-Y79AA1000976	0.66
	Y79AA1000985	F-Y79AA1000985	0.66
	Y79AA1001023	F-Y79AA1001023	0. 33
	Y79AA1001041	F-Y79AA1001041	0. 90
20	Y79AA1001048	F-Y79AA1001048	0. 81
	Y79AA1001068	F-Y79AA1001068	0.64
	Y79AA1001077	F-Y79AA1001077	0. 78
	Y79AA1001078	F-Y79AA1001078	0. 92
25	Y79AA1001105	F-Y79AA1001105	0.70
	Y79AA1001145	F-Y79AA1001145	0.46
	Y79AA1001177	F-Y79AA1001177	0. 74
	Y79AA1001185	F-Y79AA1001185	0. 74
30	Y79AA1001211	F-Y79AA1001211	0. 94
30	Y79AA1001216	F-Y79AA1001216	0. 61
	Y79AA1001228	F-Y79AA1001228	0. 56
	Y79AA1001233	F-Y79AA1001233	0. 83
	Y79AA1001236	F-Y79AA1001236	0. 32
35	Y79AA1001281	F-Y79AA1001281	0. 51
•	Y79AA1001299	F-Y79AA1001299	0. 94
	Y79AA1001312	F-Y79AA1001312	0. 53
	Y79AA1001323	F-Y79AA1001323	0. 76
40	Y79AA1001391	F~Y79AA1001391	0. 83
	Y79AA1001394	F-Y79AA1001394	0.94
	Y79AA1001402	F~Y79AA1001402	0. 90
	Y79AA1001493	F-Y79AA1001493	0. 94
45	Y79AA1001511	F-Y79AA1001511	0. 47
	Y79AA1001533	F-Y79AA1001533	0. 92
	Y79AA1001541	F-Y79AA1001541	0. 32
	Y79AA1001548	F-Y79AA1001548	0.50
50	Y79AA1001555 Y79AA1001581	F-Y79AA1001555	0. 92
		F-Y79AA1001581	0.94
	Y79AA1001585	F-Y79AA1001585	0.80
	Y79AA1001603 Y79AA1001613	F-Y79AA1001603	0.94
55		F-Y79AA1001613	0.94
55	Y79AA1001647	F-Y79AA1001647	0. 49

	Y79AA1001665	F-Y79AA1001665	0.78
	Y79AA1001679	F-Y79AA1001679	0.94
	Y79AA1001696	F-Y79AA1001696	0.60
5	Y79AA1001705	F-Y79AA1001705_	_0. 67
	Y79AA1001711	F-Y79AA1001711	0. 59
	Y79AA1001781	F-Y79AA1001781	0. 74
10	Y79AA1001805	F-Y79AA1001805	0. 79
10	Y79AA1001827	F-Y79AA1001827	0. 84
	Y79AA1001846	F-Y79AA1001846	0. 34
	Y79AA1001866	F-Y79AA1001866	0. 75
15	Y79AA1001874	F-Y79AA1001874	0. 59
	Y79AA1001923	F-Y79AA1001923	0.32
	Y79AA1001963	F-Y79AA1001963	0. 51
	Y79AA1002027	F-Y79AA1002027	
20	Y79AA1002083	F-Y79AA1002083	0. 65
	Y79AA1002089	F-Y79AA1002089	0.94
	Y79AA1002093	F-Y79AA1002093	0. 89
	Y79AA1002115	F-Y79AA1002115	0. 31
25	Y79AA1002125	F-Y79AA1002125	0. 76
	Y79AA1002139	F-Y79AA1002139	0.32
	Y79AA1002204	F-Y79AA1002204	0.94
	Y79AA1002208	F-Y79AA1002208	0.76
30	Y79AA1002209	F-Y79AA1002209	0.91
	Y79AA1002210	F-Y79AA1002210	0. 60
	Y79AA1002229	F-Y79AA1002229	0.76
35	Y79AA1002234	F-Y79AA1002234	0.94
33	Y79AA1002246	F-Y79AA1002246	0.65
	Y79AA1002258	F-Y79AA1002258	0. 79
	Y79AA1002298	F-Y79AA1002298	0.83
40	Y79AA1002307	F-Y79AA1002307	0.52
	Y79AA1002311	F-Y79AA1002311	0.92
	Y79AA1002351	F-Y79AA1002351	0.85
	Y79AA1002361	F-Y79AA1002361	0.90
45	Y79AA1002399	F-Y79AA1002399	0.94
	Y79AA1002407	F~Y79AA1002407	0.83
	Y79AA1002416	F-Y79AA1002416	0.77
	Y79AA1002431	F-Y79AA1002431	0.61
50	Y79AA1002433	F-Y79AA1002433	0.81
	Y79AA1002472	F-Y79AA1002472	0.39
	Y79AA1002482	F-Y79AA1002482	0.88
	Y79AA1002487	F-Y79AA1002487	0.87
55			

Table 10

Selected clones having the maximal ATGpr1 score of not more than 0.3 (1857 clones including the 9 clones whose 5'-end sequence does not contain the ATG codon. Blank of the maximal ATGpr1 score indicates that clones whose 5'-end sequence does not contain the ATG codon).

	clone n	ame of	maximal
	name s	equence	ATGpr1
5		-	score
	HEMBA1000042	F-HEMBA1000	042 0.08
	HEMBA1000111	F-HEMBA1000	111 0.1
10	HEMBA1000180	F-HEMBA1000	180 0.09
	HEMBA1000213	F-HEMBA1000	213 0. 21
	HEMBA1000243	F-HEMBA1000	243 0.1
	HEMBA1000251	F-HEMBA1000	251 0.08
15	HEMBA1000264	F-HEMBA1000	264 0.08
	HEMBA1000280	F-HEMBA1000	280 0.13
	HEMBA1000282	F-HEMBA1000	282 , 0. 1
	HEMBA1000333	F-HEMBA1000	333 0.17
20	HEMBA1000338	F-HEMBA1000	338 0.16
	HEMBA1000351	F-HEMBA1000	351 0.17
	HEMBA1000355	F-HEMBA1000	355 0.11
	HEMBA1000357	F-HEMBA1000	357 0.08
25	HEMBA1000366	F-HEMBA1000	366 0.11
25	HEMBA1000376	F-HEMBA1000	376 0.1
	HEMBA1000390	F-HEMBA1000	390 0.11
	HEMBA1000411	F-HEMBA1000	411 0.07
	HEMBA1000418	F-HEMBA1000	418 0.06
30	HEMBA1000422	F-HEMBA1000	422 0.07
	HEMBA1000428	F-HEMBA1000	428 0.08
	HEMBA1000434	F-HEMBA1000	434 0.11
	HEMBA1000442	F-HEMBA1000	442 0.1
35	HEMBA1000459	F-HEMBA1000	459 0.08
	HEMBA1000464	F-HEMBA1000	464 0.12
	HEMBA1000469		469 0.07
	HEMBA1000504		
40	HEMBA1000518		518 0.23
	HEMBA1000519	F-HEMBA1000	519 0.3
	HEMBA1000540		
	HEMBA1000545		
45	HEMBA1000557		
	HEMBA1000563		
•	HEMBA1000569		569 0.12
	HEMBA1000575		
50	HEMBA1000604		
	HEMBA1000622		
	HEMBA1000655		
	HEMBA1000662		
55	HEMBA1000673		
	HEMBA1000702	F-HEMBA1000	702 0.12

	HEMBA1000705	F-HEMBA1000705	0.06
	HEMBA1000722	F-HEMBA1000722	0. 17
	HEMBA1000726	F-HEMBA1000726	0.08
5	HEMBA1000747	F-HEMBA1000747	0.06
	HEMBA1000749	F-HEMBA1000749	0.09
	HEMBA1000769	F-HEMBA1000769	0. 19
	HEMBA1000773	F-HEMBA1000773	0. 08
10	HEMBA1000774	F-HEMBA1000774	0. 17
	HEMBA1000791	F-HEMBA1000791	0. 08
	HEMBA1000822	F-HEMBA1000822	0. 19
	HEMBA1000827	F-HEMBA1000827	0.09
15	HEMBA1000843	F-HEMBA1000843	0. 12
	HEMBA1000852	F-HEMBA1000852	0. 16
	HEMBA1000870	F-HEMBA1000870,	
	HEMBA1000876	F-HEMBA1000876	0. 15
20	HEMBA1000908	F-HEMBA1000908	0. 17
20	HEMBA1000934	F-HEMBA1000934	0. 2
	HEMBA1000943	F-HEMBA1000943	0.08
	HEMBA1000960	F-HEMBA1000960	0.09
25	HEMBA1000972	F-HEMBA1000972	0.12
25	HEMBA1000974	F-HEMBA1000974	0.08
	HEMBA1000985	F-HEMBA1000985	0.06
	HEMBA1000991	F-HEMBA1000991	0. 23
•	HEMBA1001007	F-HEMBA1001007	0.06
30	HEMBA1001017	F-HEMBA1001017	0.06
	HEMBA1001019	F-HEMBA1001019	0.08
	HEMBA1001020	F-HEMBA1001020	0. 15
	HEMBA1001024	F-HEMBA1001024	0.1
35	HEMBA1001026	F-HEMBA1001026	0. 13
	HEMBA1001051	F-HEMBA1001051	0. 27
	HEMBA1001060	F-HEMBA1001060	0.11
	HEMBA1001071	F-HEMBA1001071	0.1
40	HEMBA1001077	F-HEMBA1001077	0. 07
	HEMBA1001094	F-HEMBA1001094	0.07
	HEMBA1001099		0.08
	HEMBA1001121	F-HEMBA1001121	0. 25
45	HEMBA1001123	F-HEMBA1001123	0. 05
	HEMBA1001172	F-HEMBA1001172	0. 08
	HEMBA1001208	F-HEMBA1001208	0.11
	HEMBA1001226	F-HEMBA1001226	0. 27
50	HEMBA1001265	F-HEMBA1001265	0.06
	HEMBA1001294	F-HEMBA1001294	0.12
	HEMBA1001299	F-HEMBA1001299	0. 2
	HEMBA1001319	F-HEMBA1001319	0.3
55	HEMBA1001323	F-HEMBA1001323	0.07
	HEMBA1001327	F-HEMBA1001327	0. 08

	HEMBA1001330	F-HEMBA1001330	0. 11
	HEMBA1001375	F-HEMBA1001375	0. 15
	HEMBA1001383	F-HEMBA1001383	0. 11
5	HEMBA1001391	F-HEMBA1001391	
	HEMBA1001411	F-HEMBA1001411	0.06
	HEMBA1001432	F-HEMBA1001432	0. 07
	HEMBA1001433	F-HEMBA1001433	0. 24
10	HEMBA1001435	F-HEMBA1001435	0. 16
	HEMBA1001442	F-HEMBA1001442	0. 10
	HEMBA1001454	F-HEMBA1001454	0. 11
	HEMBA1001463	F-HEMBA1001463	0. 08
15	HEMBA1001478	F-HEMBA1001478	0. 08
13	HEMBA1001515	F-HEMBA1001515	0. 13
	HEMBA1001517		
	HEMBA1001517	F-HEMBA1001517,	
	HEMBA1001557	F-HEMBA1001522 F-HEMBA1001557	0.3 0.06
20	HEMBA1001566		
	HEMBA1001585	F-HEMBA1001566	0.1
	HEMBA1001589	F-HEMBA1001585 F-HEMBA1001589	0.09
	HEMBA1001608	F-HEMBA1001608	0.06
25	HEMBA1001636	F-HEMBA1001636	0.17
	HEMBA1001651		0.15
	HEMBA1001658	F-HEMBA1001651 F-HEMBA1001658	0.23
	HEMBA1001675	F-HEMBA1001675	0.17
30	HEMBA1001678		0.11
	HEMBA1001678	F-HEMBA1001678 F-HEMBA1001681	0.06
	HEMBA1001709	F-HEMBA1001709	0.1
	HEMBA1001712	F-HEMBA1001712	0. 29 0. 08
35	HEMBA1001718	F-HEMBA1001718	0. 08
	HEMBA1001734	F-HEMBA1001734	0. 07
	HEMBA1001745	F-HEMBA1001745	0. 09
	HEMBA1001761	F-HEMBA1001761	0. 09
40	HEMBA1001784	F-HEMBA1001784	0. 11
	HEMBA1001791	F-HEMBA1001791	0. 15
	HEMBA1001803	F-HEMBA1001803	0. 10
	HEMBA1001808	F-HEMBA1001808	0. 18
	HEMBA1001820	F-HEMBA1001820	0. 08
45	HEMBA1001835	F-HEMBA1001835	0. 29
	HEMBA1001844	F-HEMBA1001844	0. 11
	HEMBA1001861	F-HEMBA1001861	0. 2
	HEMBA1001888	F-HEMBA1001888	0. 18
50	HEMBA1001918	F-HEMBA1001918	0. 09
	HEMBA1001940	F-HEMBA1001940	0. 21
	HEMBA1001942	F-HEMBA1001942	0. 17
	HEMBA1001945	F-HEMBA1001945	0.06
55	HEMBA1001960	F-HEMBA1001960	-, -,

	HEMBA1001962	F-HEMBA1001962	0. 08
	HEMBA1001964	F-HEMBA1001964	0. 15
	HEMBA1001979	F-HEMBA1001979	0. 07
5	HEMBA1001991	F-HEMBA1001991	0. 08
	HEMBA1002003	F-HEMBA1002003	0. 11
	HEMBA1002008	F-HEMBA1002008	0. 08
	HEMBA1002022	F-HEMBA1002022	0. 08
10	HEMBA1002039	F-HEMBA1002039	0. 2
10	HEMBA1002100	F-HEMBA1002100	0. 07
	HEMBA1002113	F-HEMBA1002113	0. 09
	HEMBA1002139	F-HEMBA1002139	0. 2
	HEMBA1002144	F-HEMBA1002144	0. 11
15	HEMBA1002153	F-HEMBA1002153	0. 09
	HEMBA1002160	F-HEMBA1002160,	
	HEMBA1002162	F-HEMBA1002162	0. 19
	HEMBA1002166	F-HEMBA1002166	0. 06
20	HEMBA1002185	F-HEMBA1002185	0.11
	HEMBA1002204	F-HEMBA1002204	0. 09
	HEMBA1002226	F-HEMBA1002226	0. 13
	HEMBA1002253	F-HEMBA1002253	0. 05
25	HEMBA1002257	F-HEMBA1002257	0.09
	HEMBA1002270	F-HEMBA1002270	0.06
	HEMBA1002321	F-HEMBA1002321	0. 07
	HEMBA1002328	F-HEMBA1002328	0.06
30	HEMBA1002337	F-HEMBA1002337	0.11
	HEMBA1002348	F-HEMBA1002348	0.12
	HEMBA1002349	F-HEMBA1002349	0. 05
	HEMBA1002381	F-HEMBA1002381	0.09
35	HEMBA1002389	F-HEMBA1002389	0.14
	HEMBA1002486	F-HEMBA1002486	0.1
	HEMBA1002498	F-HEMBA1002498	0.17
	HEMBA1002538	F-HEMBA1002538	0. 27
40	HEMBA1002552	F-HEMBA1002552	0. 23
	HEMBA1002555	F-HEMBA1002555	0.11
	HEMBA1002558	F-HEMBA1002558	0.11
	HEMBA1002561	F-HEMBA1002561	0. 11
45	HEMBA1002590	F-HEMBA1002590	0.06
•	HEMBA1002592	F-HEMBA1002592	0. 17
	HEMBA1002621	F-HEMBA1002621	0.12
	HEMBA1002628	F-HEMBA1002628	0.06
50	HEMBA1002629	F-HEMBA1002629	0. 23
50	HEMBA1002645	F-HEMBA1002645	0.12
	HEMBA1002651	F-HEMBA1002651	0.08
	HEMBA1002659	F-HEMBA1002659	0. 28
	HEMBA1002661	F-HEMBA1002661	0. 28
55	HEMBA1002666	F-HEMBA1002666	0.16

	HEMBA1002678	F-HEMBA1002678	0. 07
	HEMBA1002679	F-HEMBA1002679	0.14
_	HEMBA1002712	F-HEMBA1002712	0. 26
5	HEMBA1002716	F-HEMBA1002716	0.14
	HEMBA1002728	F-HEMBA1002728	0.09
	HEMBA1002730	F-HEMBA1002730	0. 07
	HEMBA1002742	F-HEMBA1002742	0.09
10	HEMBA1002748	F-HEMBA1002748	0.17
	HEMBA1002780	F-HEMBA1002780	0.1
	HEMBA1002801	F-HEMBA1002801	0.1
	HEMBA1002826	F-HEMBA1002826	0. 1
15	HEMBA1002833	F-HEMBA1002833	0.14
	HEMBA1002886	F-HEMBA1002886	0.09
	HEMBA1002896	F-HEMBA1002896,	0.05
	HEMBA1002921	F-HEMBA1002921	0. 25
20	HEMBA1002924	F-HEMBA1002924	0.12
	HEMBA1002934	F-HEMBA1002934	0. 2
	HEMBA1002944	F-HEMBA1002944	0. 12
	HEMBA1002968	F-HEMBA1002968	0.12
25	HEMBA1003034	F-HEMBA1003034	0.06
	HEMBA1003037	F-HEMBA1003037	0.11
	HEMBA1003064	F-HEMBA1003064	0.09
	HEMBA1003071	F-HEMBA1003071	0. 27
30	HEMBA1003083	F-HEMBA1003083	0. 18
	HEMBA1003086	F-HEMBA1003086	0.06
	HEMBA1003098	F-HEMBA1003098	0.09
	HEMBA1003133	F-HEMBA1003133	0. 12
35	HEMBA1003142	F-HEMBA1003142	0. 29
33	HEMBA1003166	F-HEMBA1003166	0. 1
	HEMBA1003197	F-HEMBA1003197	0.1
	HEMBA1003202 HEMBA1003204	F-HEMBA1003202 F-HEMBA1003204	0.17
40	HEMBA1003212	F-HEMBA1003212	0. 13 0. 13
40	HEMBA1003212	F-HEMBA1003212	
	HEMBA1003229		0. 14
	HEMBA1003273	F-HEMBA1003229 F-HEMBA1003273	0. 19 0. 1
	HEMBA1003276	F-HEMBA1003276	0. 12
45	HEMBA1003278	F-HEMBA1003278	0. 12
	HEMBA1003296	F-HEMBA1003296	0. 03
	HEMBA1003304	F-HEMBA1003304	0.09
	HEMBA1003304	F-HEMBA1003309	0. 09
50	HEMBA1003314	F-HEMBA1003314	0. 07
	HEMBA1003328	F-HEMBA1003328	0. 13
	HEMBA1003330	F-HEMBA1003330	0. 13
	HEMBA1003348	F-HEMBA1003348	0. 17
55	HEMBA1003373	F-HEMBA1003373	0. 21

	HEMBA1003376	F-HEMBA1003376	0.14
	HEMBA1003384	F-HEMBA1003384	0.07
	HEMBA1003531	F-HEMBA1003531	0. 2
5	HEMBA1003548	F-HEMBA1003548	0.08
	HEMBA1003571	F-HEMBA1003571	0, 13
	HEMBA1003579	F-HEMBA1003579	0.13
	HEMBA1003595	F-HEMBA1003595	0.08
10	HEMBA1003597	F-HEMBA1003597	0.1
	HEMBA1003598	F-HEMBA1003598	0.12
	HEMBA1003622	F-HEMBA1003622	0.08
	HEMBA1003630	F-HEMBA1003630	0.08
15	HEMBA1003637	F-HEMBA1003637	0.07
	HEMBA1003640	F-HEMBA1003640	0.16
	HEMBA1003656	F-HEMBA1003656,	0.13
	HEMBA1003714	F-HEMBA1003714	0.12
20	HEMBA1003715	F-HEMBA1003715	0.1
	HEMBA1003725	F-HEMBA1003725	0.08
	HEMBA1003733	F-HEMBA1003733	0. 2
	HEMBA1003758	F-HEMBA1003758	0.06
25	HEMBA1003773	F-HEMBA1003773	0.11
-	HEMBA1003784	F-HEMBA1003784	0.09
	HEMBA1003856	F-HEMBA1003856	0. 25
	HEMBA1003885	F-HEMBA1003885	0.09
30	HEMBA1003902	F-HEMBA1003902	0.08
30	HEMBA1003908	F-HEMBA1003908	0.08
	HEMBA1003926	F-HEMBA1003926	0.09
	HEMBA1003937	F-HEMBA1003937	0.2
35	HEMBA1003939	F-HEMBA1003939	0. 12
35	HEMBA1003942 HEMBA1003950	F-HEMBA1003942 F-HEMBA1003950	0.1
	HEMBA1003958	F-HEMBA1003958	0. 09 0. 07
	HEMBA1003976	F-HEMBA1003976	0. 07
40	HEMBA1003987	F-HEMBA1003987	0. 24
40	HEMBA1004000		0.16
	HEMBA1004012	F-HEMBA1004012	0. 12
	HEMBA1004015	F-HEMBA1004015	0. 23
	HEMBA1004024	F-HEMBA1004024	0.13
45	HEMBA1004038	F-HEMBA1004038	0.09
	HEMBA1004042	F-HEMBA1004042	0.09
	HEMBA1004045	F-HEMBA1004045	0.09
	HEMBA1004049	F-HEMBA1004049	0. 09
50	HEMBA1004132	F-HEMBA1004132	0.08
	HEMBA1004138	F-HEMBA1004138	0.07
	HEMBA1004164	F-HEMBA1004164	0.08
	HEMBA1004225	F-HEMBA1004225	0.1
55	HEMBA1004241	F-HEMBA1004241	0.08

	HEMBA1004267	F-HEMBA1004267	0.09
	HEMBA1004272	F-HEMBA1004272	0.09
	HEMBA1004295	F-HEMBA1004295	0.19
5	HEMBA1004306	F-HEMBA1004306	0. 07
	HEMBA1004312	F-HEMBA1004312	0. 26
	HEMBA1004323	F-HEMBA1004323	0. 13
	HEMBA1004334	F-HEMBA1004334	0.06
10	HEMBA1004354	F-HEMBA1004354	0.1
	HEMBA1004356	F-HEMBA1004356	0.07
	HEMBA1004366	F-HEMBA1004366	0.14
	HEMBA1004394	F-HEMBA1004394	0. 08
15	HEMBA1004396	F-HEMBA1004396	0.05
10	HEMBA1004405	F-HEMBA1004405	0. 17
	HEMBA1004429	F-HEMBA1004429,	
	HEMBA1004433	F-HEMBA1004433	0. 29
•	HEMBA1004460	F-HEMBA1004460	0.11
20	HEMBA1004461	F-HEMBA1004461	0.11
	HEMBA1004482	F-HEMBA1004482	0.15
	HEMBA1004506	F-HEMBA1004506	0.05
	HEMBA1004538	F-HEMBA1004538	0.15
25	HEMBA1004577	F-HEMBA1004577	0.17
	HEMBA1004586	F-HEMBA1004586	0.06
	HEMBA1004617	F-HEMBA1004617	0.11
	HEMBA1004629	F-HEMBA1004629	0:11
30	HEMBA1004631	F-HEMBA1004631	0.1
	HEMBA1004666	F-HEMBA1004666	0.08
	HEMBA1004670	F-HEMBA1004670	0.15
	HEMBA1004672	F-HEMBA1004672	0.14
35	HEMBA1004730	F-HEMBA1004730	0. 15
	HEMBA1004733	F-HEMBA1004733	0.12
	HEMBA1004748	F-HEMBA1004748	0.1
	HEMBA1004770	F-HEMBA1004770	0. 15
40	HEMBA1004778	F-HEMBA1004778	0.14
	HEMBA1004803	F-HEMBA1004803	0.13
	HEMBA1004807	F-HEMBA1004807	0.18
	HEMBA1004816	F-HEMBA1004816	0.06
45	HEMBA1004820	F-HEMBA1004820	0.11
	HEMBA1004865	F-HEMBA1004865	0. 17
	HEMBA1004880	F-HEMBA1004880	0. 28
	HEMBA1004900	F-HEMBA1004900	0.11
50	HEMBA1004909	F-HEMBA1004909	0. 11
	HEMBA1004918	F-HEMBA1004918	0.09
	HEMBA1004956	F-HEMBA1004956	0. 05
	HEMBA1004960	F-HEMBA1004960	0. 21
	HEMBA1004978	F-HEMBA1004978	0. 1
55	HEMBA1004983	F-HEMBA1004983	0. 28

	HEMBA1004995	F-HEMBA1004995	0. 24
	HEMBA1005008	F-HEMBA1005008	0. 07
	HEMBA1005039	F-HEMBA1005039	0. 08
5	HEMBA1005062	F-HEMBA1005062	0.11
	HEMBA1005123	F-HEMBA1005123	0. 07
	HEMBA1005152	F-HEMBA1005152	0.11
	HEMBA1005159	F-HEMBA1005159	0.1
10	HEMBA1005232	F-HEMBA1005232	0.1
	HEMBA1005241	F-HEMBA1005241	0.14
	HEMBA1005251	F-HEMBA1005251	0. 29
	HEMBA1005274	F-HEMBA1005274	0.06
15	HEMBA1005275	F-HEMBA1005275	0.12
	HEMBA1005293	F-HEMBA1005293	0.06
	HEMBA1005304	F-HEMBA1005304,	0.09
	HEMBA1005311	F-HEMBA1005311	0. 2
20	HEMBA1005315	F-HEMBA1005315	0.07
	HEMBA1005318	F-HEMBA1005318	0.08
	HEMBA1005353	F-HEMBA1005353	0.12
	HEMBA1005359	F-HEMBA1005359	0. 15
25	HEMBA1005372	F-HEMBA1005372	0. 07
20	HEMBA1005374	F-HEMBA1005374	0.09
	HEMBA1005389	F-HEMBA1005389	0.1
	HEMBA1005408	F-HEMBA1005408	0.09
20	HEMBA1005410	F-HEMBA1005410	0. 07
30	HEMBA1005426	F-HEMBA1005426	0.16
	HEMBA1005443	F-HEMBA1005443	0. 17
	HEMBA1005447	F-HEMBA1005447	0. 25
	HEMBA1005497	F-HEMBA1005497	0.09
35	HEMBA1005500	F-HEMBA1005500	0.08
	HEMBA1005506	F-HEMBA1005506	0.17
	HEMBA1005508	F-HEMBA1005508	0.12
	HEMBA1005511	F-HEMBA1005511	0.1
40	HEMBA1005520	F-HEMBA1005520	0.07
	HEMBA1005552	F-HEMBA1005552	0. 05
	HEMBA1005568	F-HEMBA1005568	0.06
	HEMBA1005570	F-HEMBA1005570	0.1
45	HEMBA1005577	F-HEMBA1005577	0.13
	HEMBA1005588	F-HEMBA1005588	0. 25
	HEMBA1005593	F-HEMBA1005593	0.13
	HEMBA1005606 HEMBA1005616	F-HEMBA1005606	0.16
50	HEMBA1005627	F-HEMBA1005616	0. 15
	HEMBA1005627	F-HEMBA1005627 F-HEMBA1005631	0.11
	HEMBA1005632	F-HEMBA1005632	0.07
	HEMBA1005634	F-HEMBA1005634	0. 07 0. 07
55	HEMBA1005670	F-HEMBA1005670	
	וובשטא ו 10000 /0	L-UEWDW 1009010	0.11

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	HEMBA1005679	F~HEMBA1005679	0. 22
	HEMBA1005699	F-HEMBA1005699	0.08
_	HEMBA1005705	F-HEMBA1005705	0.08
5	HEMBA1005717	F-HEMBA1005717	0.17
	HEMBA1005765	F-HEMBA1005765	0.08
	HEMBA1005780	F-HEMBA1005780	0.16
	HEMBA1005829	F-HEMBA1005829	0.09
10	HEMBA1005853	F-HEMBA1005853	0.06
	HEMBA1005894	F-HEMBA1005894	0.16
	HEMBA1005909	F-HEMBA1005909	0.13
	HEMBA1005911	F-HEMBA1005911	0. 22
15	HEMBA1005921	F-HEMBA1005921	0.1
	HEMBA1005931	F-HEMBA1005931	0.16
	HEMBA1005934	F-HEMBA1005934,	0. 1
	HEMBA1005962	F-HEMBA1005962	0.07
20	HEMBA1005999	F-HEMBA1005999	0.14
20	HEMBA1006002	F-HEMBA1006002	0.09
	HEMBA1006005	F-HEMBA1006005	0. 2
	HEMBA1006036	F-HEMBA1006036	0. 24
	HEMBA1006042	F-HEMBA1006042	0.11
25	HEMBA1006067	F-HEMBA1006067	0.09
	HEMBA1006090	F-HEMBA1006090	0.09
	HEMBA1006124	F-HEMBA1006124	0.05
	HEMBA1006142	F-HEMBA1006142	0.13
30	HEMBA1006158	F-HEMBA1006158	0. 27
	HEMBA1006253	F-HEMBA1006253	0. 29
	HEMBA1006259	F-HEMBA1006259	0.11
	HEMBA1006268	F-HEMBA1006268	0.11
35	HEMBA1006328	F-HEMBA1006328	0.06
	HEMBA1006334	F-HEMBA1006334	0.08
	HEMBA1006359	F-HEMBA1006359	0. 22
	HEMBA1006364	F-HEMBA1006364	0. 23
40	HEMBA1006380	F-HEMBA1006380	0.09
	HEMBA1006416	F-HEMBA1006416	0. 25
	HEMBA1006419	F-HEMBA1006419	0.08
	HEMBA1006421	F-HEMBA1006421	0.26
45	HEMBA1006426	F-HEMBA1006426	0.17
•	HEMBA1006438	F-HEMBA1006438	0.08
	HEMBA1006446	F-HEMBA1006446	0.11
	HEMBA1006461	F-HEMBA1006461	0.15
50	HEMBA1006471	F-HEMBA1006471	0.06
50	HEMBA1006486	F-HEMBA1006486	0.11
	HEMBA1006489	F-HEMBA1006489	0.09
	HEMBA1006540	F-HEMBA1006540	0. 23
	HEMBA1006546	F-HEMBA1006546	0.07
55	HEMBA1006562	F-HEMBA1006562	0.12

	HEMBA1006569	F-HEMBA1006569	
	HEMBA1006595	F~HEMBA1006595	0.08
	HEMBA1006597	F-HEMBA1006597	0.17
5	HEMBA1006617	F-HEMBA1006617	
	HEMBA1006635	F-HEMBA1006635	0. 07
	HEMBA1006639	F-HEMBA1006639	0. 25
	HEMBA1006648	F-HEMBA1006648	0. 24
10	HEMBA1006653	F-HEMBA1006653	0.09
	HEMBA1006665	F-HEMBA1006665	0. 17
	HEMBA1006676	F-HEMBA1006676	0. 17
	HEMBA1006695	F-HEMBA1006695	0.1
15	HEMBA1006696	F-HEMBA1006696	0.05
	HEMBA1006744	F-HEMBA1006744	0, 12
	HEMBA1006779	F-HEMBA1006779,	0. 07
	HEMBA1006780	F-HEMBA1006780	0. 23
20	HEMBA1006821	F~HEMBA1006821	0. 08
	HEMBA1006824	F-HEMBA1006824	0.09
	HEMBA1006849	F-HEMBA1006849	0.05
	HEMBA1006865	F-HEMBA1006865	0. 1
25	HEMBA1006921	F-HEMBA1006921	0. 22
25	HEMBA1006929	F-HEMBA1006929	0. 15
	HEMBA1006938	F-HEMBA1006938	0.07
	HEMBA1006949	F-HEMBA1006949	0. 27
	HEMBA1007017	F-HEMBA1007017	0.07
30	HEMBA1007045	F-HEMBA1007045	0.14
	HEMBA1007051	F-HEMBA1007051	0. 19
	HEMBA1007073	F-HEMBA1007073	0. 22
	HEMBA1007078	F-HEMBA1007078	0.12
35	HEMBA1007113	F-HEMBA1007113	0.12
	HEMBA1007129	F-HEMBA1007129	0. 18
	HEMBA1007147	F-HEMBA1007147	0.08
	HEMBA1007206	F-HEMBA1007206	0. 11
40	HEMBA1007256	F-HEMBA1007256	0.13
	HEMBA1007273	F-HEMBA1007273	0.05
	HEMBA1007279	F-HEMBA1007279	0.09
	HEMBA1007288	F-HEMBA1007288	0.06
45	HEMBA1007322	F-HEMBA1007322	0. 12
	HEMBA1007327	F-HEMBA1007327	0.12
	HEMBA1007341	F-HEMBA1007341	0.1
	HEMBA1007347	F-HEMBA1007347	0.14
50	HEMBB1000005	F-HEMBB1000005	0. 13
	HEMBB1000039	F-HEMBB1000039	0.07
	HEMBB1000044	F-HEMBB1000044	0.08
	HEMBB1000050	F-HEMBB1000050	0.07
55	HEMBB1000054	F-HEMBB1000054	0.12
	HEMBB1000055	F-HEMB81000055	0. 2

	HEMBB1000059	F-HEMBB1000059	0. 08
	HEMBB1000089	F-HEMBB1000089	0.09
	HEMBB1000099	F-HEMBB1000099	0. 17
5	HEMBB1000113	F-HEMBB1000113	0.12
	HEMBB1000141	F-HEMBB1000141	0.13
	HEMBB1000144	F-HEMBB1000144	0.2
	HEMBB1000218	F-HEMBB1000218	0.07
10	HEMBB1000250	F-HEMBB1000250	0.26
	HEMB81000258	F-HEMBB1000258	0. 15
	HEMBB1000264	F-HEMBB1000264	0.08
	HEMBB1000272	F-HEMBB1000272	0.11
15	HEMBB1000274	F-HEMBB1000274	0.12
	HEMBB1000284	F-HEMBB1000284	0.1
	HEMBB1000307	F-HEMBB1000307,	0.11
	HEMBB1000312	F-HEMBB1000312	0.07
20	HEMBB1000317	F-HEMBB1000317	0. 27
20	HEMBB1000318	F-HEMBB1000318	0.11
	HEMBB1000335	F-HEMBB1000335	0. 15
	HEMBB1000336	F-HEMBB1000336	0. 28
	HEMBB1000337	F-HEMBB1000337	0. 25
25	HEMBB1000341	F-HEMBB1000341	0. 24
	HEMBB1000343	F-HEMBB1000343	0. 07
	HEMBB1000354	F-HEMBB1000354	0. 15
	HEMBB1000369	F-HEMBB1000369	0.08
30	HEMBB1000374	F-HEMBB1000374	0. 3
	HEMBB1000376	F-HEMBB1000376	0.12
	HEMBB1000399	F-HEMBB1000399	0.08
	HEMBB1000402	F-HEMBB1000402	0. 29
35	HEMBB1000404	F-HEMBB1000404	0. 29
	HEMBB1000420	F-HEMBB1000420	0.11
	HEMBB1000434	F-HEMBB1000434	0.05
	HEMBB1000441	F-HEMBB1000441	0.12
40	HEMBB1000455	F-HEMBB1000455	0.05
	HEMBB1000472	F-HEMBB1000472	0.05
	HEMBB1000480	F-HEMBB1000480	0. 13
	HEMBB1000487	F-HEMBB1000487	0.09
45	HEMBB1000490	F-HEMBB1000490	0.06
	HEMBB1000491	F-HEMBB1000491	0.16
	HEMBB1000493	F-HEMBB1000493	0. 15
	HEMBB1000518	F-HEMBB1000518	0.05
50	HEMBB1000523	F-HEMBB1000523	0.07
	HEMBB1000530	F-HEMBB1000530	0.14
	HEMBB1000554	F-HEMBB1000554	0. 23
	HEMBB1000564	F-HEMBB1000564	0.09
55	HEMBB1000573	F-HEMBB1000573	0.1
	HEMBB1000575	F-HEMBB1000575	0. 08

	HEMBB1000586	F-HEMBB1000586	0.12
	HEMBB1000598	F-HEMBB1000598	0.12
	HEMBB1000637	F-HEMBB1000637	0.19
5	HEMBB1000638	F-HEMBB1000638	0.07
	HEMBB1000643	F-HEMBB1000643	0.07
	HEMBB1000649	F-HEMBB1000649	0.13
	HEMBB1000652	F-HEMBB1000652	0. 2
10	HEMBB1000665	F-HEMBB1000665	0.07
	HEMBB1000684	F-HEMBB1000684	0.06
	HEMBB1000709	F-HEMBB1000709	0.09
	HEMBB1000726	F-HEMBB1000726	0. 07
15	HEMBB1000738	F-HEMBB1000738	0.19
	HEMBB1000749	F-HEMBB1000749	0.14
	HEMBB1000770	F-HEMBB1000770,	0.09
	HEMBB1000790	F-HEMBB1000790	0. 23
20	HEMBB1000794	F-HEMBB1000794	0.09
20	HEMBB1000821	F-HEMBB1000821	0. 1
	HEMBB1000822	F-HEMBB1000822	0.18
	HEMBB1000827	F-HEMBB1000827	0. 1
25	HEMBB1000831	F-HEMBB1000831	0. 2
25	HEMBB1000840	F-HEMBB1000840	0.09
	HEMBB1000876	F-HEMBB1000876	0.12
	HEMBB1000883	F-HEMBB1000883	0.13
	HEMBB1000888	F-HEMBB1000888	0. 1
30	HEMBB1000890	F-HEMBB1000890	0.09
	HEMBB1000893	F-HEMBB1000893	0. 1
	HEMBB1000910	F-HEMBB1000910	0.06
	HEMBB1000913	F-HEMBB1000913	0.09
35	HEMBB1000915	F-HEMBB1000915	0. 12
	HEMBB1000917	F-HEMBB1000917	0. 15
	HEMBB1000959	F-HEMBB1000959	0.08
	HEMBB1000981	F-HEMBB1000981	0.09
40	HEMBB1000996	F-HEMBB1000996	0.1
	HEMBB1001004	F-HEMBB1001004	0.08
	HEMBB1001008	F-HEMBB1001008	0. 25
	HEMBB1001037	F-HEMBB1001037	0.14
45	HEMBB1001047	F-HEMBB1001047	0.14
	HEMBB1001051	F-HEMBB1001051	0.13
	HEMBB1001060	F-HEMBB1001060	0.16
	HEMBB1001063	F-HEMBB1001063	0.1
50	HEMB81001102	F-HEMBB1001102	0.19
	HEMB81001114	F-HEMBB1001114	0.08
	HEMBB1001119	F-HEMBB1001119	0.13
	HEMBB1001133	F-HEMB81001133	0.06
55	HEMBB1001142	F-HEMB81001142	0.17
	HEMBB1001177	F-HEMBB1001177	0. 2

	HEMBB1001208	F-HEMBB1001208	0. 12
	HEMBB1001209	F-HEMBB1001209	0.11
	HEMBB1001218	F-HEMB81001218	0.1
5	HEMB81001221	F-HEMBB1001221	0.1
	HEMBB1001249	F-HEMBB1001249	0.12
	HEMBB1001253	F-HEMBB1001253	0.13
	HEMBB1001254	F-HEMBB1001254	0.13
10	HEMBB1001267	F-HEMBB1001267	0.09
	HEMBB1001271	F-HEMBB1001271	0.16
	HEMBB1001282	F-HEMBB1001282	0.14
	HEMBB1001302	F-HEMBB1001302	0. 18
15	HEMBB1001304	F-HEMBB1001304	0.17
	HEMBB1001315	F-HEMBB1001315	0.18
	HEMBB1001317	F-HEMBB1001317,	0.12
	HEMBB1001326	F-HEMBB1001326	0.09
20	HEMBB1001335	F-HEMBB1001335	0.08
20	HEMBB1001337	F-HEMBB1001337	0. 12
	HEMBB1001348	F-HEMBB1001348	0.11
	HEMBB1001356	F-HEMBB1001356	0.1
0.5	HEMBB1001366	F-HEMBB1001366	0.16
25	HEMBB1001367	F-HEMB81001367	0.11
	HEMBB1001380	F-HEMBB1001380	0. 07
	HEMBB1001424	F-HEMBB1001424	0. 05
	HEMBB1001436	F-HEMBB1001436	0.14
30	HEMBB1001443	F-HEMBB1001443	0. 28
	HEMBB1001454	F-HEMBB1001454	
	HEMBB1001458	F-HEMBB1001458	0.11
	HEMBB1001463	F-HEMBB1001463	0. 1
35	HEMBB1001464	F-HEMBB1001464	0.06
	HEMBB1001500	F-HEMBB1001500	0. 11
	HEMBB1001521	F-HEMBB1001521	0. 13
	HEMBB1001527	F-HEMBB1001527	0.13
40	HEMBB1001535	F-HEMBB1001535	0.11
	HEMBB1001536	F-HEMBB1001536	0.08
	HEMBB1001537	F-HEMBB1001537	0.11
	HEMBB1001555	F-HEMBB1001555	0.08
45	HEMBB1001565	F-HEMBB1001565	0.1
	HEMBB1001586	F-HEMBB1001586	0.06
	HEMBB1001588	F-HEMB81001588	0.06
	HEMBB1001618	F-HEMBB1001618	0.09
50	HEMB81001619	F-HEMBB1001619	0.12
	HEMB81001630	F-HEMBB1001630	0. 12
	HEMBB1001637	F-HEMBB1001637	0.08
	HEMBB1001641	F-HEMBB1001641	0.09
55	HEMBB1001704	F-HEMBB1001704	0.09
	HEMBB1001706	F-HEMBB1001706	0. 11

	HEMBB1001717	F-HEMBB1001717	0.09
	HEMBB1001735	F-HEMBB1001735	0. 21
	HEMBB1001747	F-HEMBB1001747	0.07
5	HEMBB1001749	F-HEMBB1001749	0.13
	HEMBB1001753	F-HEMBB1001753	0.08
	HEMBB1001756	F-HEMBB1001756	0.11
	HEMBB1001762	F-HEMBB1001762	0.11
10	HEMBB1001797	F-HEMBB1001797	0.13
	HEMBB1001802	F-HEMBB1001802	0.06
	HEMBB1001836	F-HEMBB1001836	0.15
	HEMBB1001850	F-HEMBB1001850	0.06
15	HEMBB1001863	F-HEMBB1001863	0.09
	HEMBB1001867	F-HEMBB1001867	0.11
	HEMBB1001868	F-HEMBB1001868,	
	HEMBB1001875	F-HEMBB1001875	0. 05
20	HEMBB1001880	F-HEMBB1001880	0.08
20	HEMBB1001899	F-HEMBB1001899	0.16
	HEMBB1001911	F-HEMBB1001911	0. 22
	HEMBB1001921	F-HEMBB1001921	0.07
	HEMBB1001922	F-HEMBB1001922	0.11
25	HEMBB1001925	F-HEMBB1001925	0.19
	HEMBB1001930	F-HEMBB1001930	0.1
	HEMBB1001944	F-HEMBB1001944	0.14
	HEMBB1001952	F-HEMBB1001952	0.08
30	HEMBB1001953	F-HEMBB1001953	0. 15
	HEMBB1001967	F-HEMBB1001967	0.08
	HEMBB1001973	F-HEMBB1001973	0.09
	HEMBB1001983	F-HEMBB1001983	0.1
35	HEMBB1001988	F-HEMBB1001988	0.15
	HEMBB1001996	F-HEMBB1001996	0.19
	HEMBB1001997	F-HEMBB1001997	0.08
	HEMBB1002002	F-HEMBB1002002	0. 21
40	HEMBB1002005	F-HEMBB1002005	0.18
	HEMBB1002009	F-HEMBB1002009	0.09
	HEMBB1002015	F-HEMBB1002015	0. 26
	HEMBB1002042	F-HEMBB1002042	0.11
45	HEMBB1002043	F-HEMBB1002043	0.09
	HEMBB1002045	F-HEMBB1002045	0. 26
	HEMBB1002049	F-HEMBB1002049	0.12
	HEMBB1002069	F-HEMBB1002069	0.12
50	HEMBB1002092	F-HEMBB1002092	0.16
	HEMBB1002094	F-HEMBB1002094	0.11
	HEMBB1002115	F-HEMBB1002115	0. 09
	HEMBB1002139	F-HEMBB1002139	0.13
55	HEMBB1002189	F-HEMBB1002189	0.06
55	HEMBB1002190	F-HEMBB1002190	0.18

	HEMBR1002232	F-HEMBB1002232	0. 09
	HEMBB1002247	F-HEMBB1002247	0. 17
	HEMBB1002254	F-HEMBB1002254	0. 17
5	HEMBB1002255		
	- ···•-	F-HEMBB1002255	• •
	HEMBB1002266	F-HEMBB1002266	0. 28
	HEMBB1002280	F-HEMBB1002280	0.07
	HEMBB1002306	F-HEMBB1002306	0.06
10	HEMBB1002364	F-HEMBB1002364	0. 05
	HEMBB1002371	F-HEMBB1002371	0. 12
	HEMBB1002381	F-HEMBB1002381	0. 09
	HEMBB1002387	F-HEMBB1002387	0. 17
15	HEMBB1002425	F-HEMB81002425	0.08
	HEMBB1002453	F-HEMBB1002453	0. 12
	HEMBB1002458	F-HEMBB1002458,	0.09
	HEMBB1002477	F-HEMBB1002477	0.17
20	HEMBB1002509	F-HEMBB1002509	0.06
	HEMBB1002520	F-HEMBB1002520	0.08
	HEMBB1002522	F-HEMBB1002522	0. 21
	HEMBB1002531	F-HEMBB1002531	0.1
25	HEMBB1002534	F-HEMBB1002534	0.09
25	HEMBB1002556	F-HEMBB1002556	0.1
	HEMBB1002579	F-HEMBB1002579	0.13
	HEMBB1002582	F-HEMBB1002582	0. 17
	HEMBB1002590	F-HEMBB1002590	0.08
30	HEMBB1002596	F-HEMBB1002596	0. 15
	HEMBB1002601	F-HEMBB1002601	0.1
	HEMBB1002603	F-HEMBB1002603	0.12
	HEMBB1002610	F-HEMBB1002610	0.1
35	HEMBB1002613	F-HEMBB1002613	0.09
	HEMBB1002617	F-HEMBB1002617	0.12
	HEMBB1002623	F-HEMBB1002623	0.14
	HEMBB1002635	F-HEMBB1002635	0.08
40	HEMBB1002664	F-HEMBB1002664	0.06
	HEMBB1002677	F-HEMBB1002677	0.07
	HEMBB1002683	F-HEMBB1002683	0.06
	HEMBB1002686	F-HEMBB1002686	0.12
45	HEMB81002699	F-HEMBB1002699	0.15
	HEMBB1002702	F-HEMBB1002702	0.16
	HEMBB1002712	F-HEMBB1002712	0.07
	MAMMA1000009	F-MAMMA1000009	0.11
	MAMMA1000043	F-MAMMA1000043	0.19
50	MAMMA1000045	F-MAMMA1000045	0.08
	MAMMA1000092	F-MAMMA1000092	0.11
	MAMMA1000103	F-MAMMA1000103	0.09
	MAMMA1000117	F-MAMMA1000117	0. 2
55	MAMMA1000129	F-MAMMA1000129	0.17

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	MAMMA1000133	F-MAMMA1000133	0. 1
	MAMMA1000134	F-MAMMA1000134	0. 05
_	MAMMA1000155	F-MAMMA1000155	0. 12
5	MAMMA1000175	F-MAMMA1000175	0. 07
	MAMMA1000198	F-MAMMA1000198	0. 2
	MAMMA1000221	F-MAMMA1000221	0. 1
	MAMMA1000227	F-MAMMA1000227	0.08
10	MAMMA1000241	F-MAMMA1000241	0.16
	MAMMA1000254	F-MAMMA1000254	0.08
	MAMMA1000264	F-MAMMA1000264	0.1
	MAMMA1000266	F-MAMMA1000266	0.18
15	MAMMA1000270	F-MAMMA1000270	0. 1
	MAMMA1000287	F-MAMMA1000287	0.1
	MAMMA1000302	F-MAMMA1000302,	0.07
	MAMMA1000307	F-MAMMA1000307	0. 26
20	MAMMA1000331	F-MAMMA1000331	0.18
	MAMMA1000340	F-MAMMA1000340	0. 2
	MAMMA1000348	F-MAMMA1000348	0.09
	MAMMA1000356	F-MAMMA1000356	0. 15
05	MAMMA1000360	F-MAMMA1000360	0.09
25	MAMMA1000385	F-MAMMA1000385	0.08
	MAMMA1000402	F-MAMMA1000402	0.11
	MAMMA1000413	F-MAMMA1000413	0.1
	MAMMA1000414	F-MAMMA1000414	0.1
30	MAMMA1000423	F-MAMMA1000423	0.06
	MAMMA1000424	F-MAMMA1000424	0. 1
	MAMMA1000431	F-MAMMA1000431	0. 28
	MAMMA1000444	F-MANMA1000444	0. 07
35	MAMMA1000446	F-MAMMA1000446	0.08
	MAMMA1000478	F-MAMMA1000478	0. 1
	MAMMA1000483	F-MAMMA1000483	0. 07
	MAMMA1000500	F-MAMMA1000500	0.16
40	MAMMA1000501	F-MAMMA1000501	0.05
	MAMMA1000516	F-MAMMA1000516	0.06
	MAMMA1000522	F-MAMMA1000522	0. 21
	MAMMA1000559	F-MAMMA1000559	0.12
45	MAMMA1000565	F-MAMMA1000565	0.09
	MAMMA1000576	F-MAMMA1000576	0.14
	MAMMA1000585	F-MAMMA1000585	0. 07
	MAMMA1000594	F-MAMMA1000594	0. 17
50	MAMMA1000597	F-MAMMA1000597	0. 1
50	MAMMA1000605	F-MAMMA1000605	0.09
•	MAMMA1000616	F-MAMMA1000616	0.08
	MAMMA1000621	F-MAMMA1000621	0. 07
	MAMMA1000643	F-MAMMA1000643	0. 1
55	MAMMA1000669	F-MAMMA1000669	0.11

	MAMMA1000696	F-MAMMA1000696	0.08
	MAMMA1000707	F-MAMMA1000707	0.07
	MAMMA1000714	F-MAMMA1000714	0.18
5	MAMMA1000718	F-MAMMA1000718	0.06
	MAMMA1000720	F-MAMMA1000720	0.15
	MAMMA1000723	F-MAMMA1000723	0.14
	MAMMA1000732	F-MAMMA1000732	0.06
10	MAMMA1000733	F-MAMMA1000733	0.1
	MAMMA1000744	F-MAMMA1000744	0.15
	MAMMA1000752	F-MAMMA1000752	0.12
	MAMMA1000761	F-MAMMA1000761	0.24
15	MAMMA1000775	F-MAMMA1000775	0.24
	MAMMA1000778	F-MAMMA1000778	0.09
	MAMMA1000782	F-MAMMA1000782,	0.1
	MAMMA1000798	F-MAMMA1000798	0.16
20	MAMMA1000802	F-MAMMA1000802	0.11
	MAMMA1000839	F-MAMMA1000839	0.09
	MAMMA1000845	F-MAMMA1000845	0. 1
	MAMMA1000851	F-MAMMA1000851	0.08
25	MAMMA1000855	F-MAMMA1000855	0. 23
	MAMMA1000862	F-MAMMA1000862	0.12
	MAMMA1000863	F-MAMMA1000863	0. 16
	MAMMA1000867	F-MAMMA1000867	0. 25
30	MAMMA1000876	F-MAMMA1000876	0.14
	MAMMA1000877	F-MAMMA1000877	0.07
	MAMMA1000880 MAMMA1000883	F-MAMMA1000880 F-MAMMA1000883	0. 13 0. 3
	MAMMA1000883	F-MAMMA1000883	0. 3
35	MAMMA1000931	F-MAMMA1000931	0. 08
	MAMMA1000940	F-MAMMA1000940	0. 18
	MAMMA1000941	F-MAMMA1000941	0. 13
	MAMMA1000942	F-MAMMA1000942	0.08
40	MAMMA1000943	F-MAMMA1000943	0.09
40	MAMMA1000957	F-MAMMA1000957	0.13
	MAMMA1000962	F-MAMMA1000962	0.06
	MAMMA1000975	F-MAMMA1000975	0. 26
45	MAMMA1000987	F-MAMMA1000987	0.08
45	MAMMA1000998	F-MAMMA1000998	0.08
	MAMMA1001003	F-MAMMA1001003	0. 15
	MAMMA1001024	F-MAMMA1001024	0.08
50	MAMMA1001030	F-MAMMA1001030	0. 2
50	MAMMA1001035	F-MAMMA1001035	0.08
	MAMMA1001038	F-MAMMA1001038	0. 15
	MAMMA1001050	F-MAMMA1001050	0.06
	MAMMA1001067	F-MAMMA1001067	0. 08
55	MAMMA1001074	F-MAMMA1001074	0. 09

	MAMMA1001082	F-MAMMA1001082	0. 1
	MAMMA1001092	F-MAMMA1001092	0. 05
	MAMMA1001032	F-MAMMA1001133	0. 03
5	MAMMA1001145		
3	MAMMA1001145	F-MAMMA1001145_ F-MAMMA1001161	0.11
			0.11
	MAMMA1001162	F-MAMMA1001162	
	MAMMA1001186	F-MAMMA1001186	0.09
10	MAMMA1001191	F-MAMMA1001191	0.08
	MAMMA1001203	F-MAMMA1001203	0.08
	MAMMA1001206	F-MAMMA1001206	0.09
	MAMMA1001220	F-MAMMA1001220	0.1
15	MAMMA1001243	F-MAMMA1001243	0.14
	MAMMA1001249	F-MAMMA1001249	0. 13
	MAMMA1001256	F-MAMMA1001256,	0. 1
	MAMMA1001268	F-MAMMA1001268	0. 26
20	MAMMA1001274	F-MAMMA1001274	0.13
	MAMMA1001280	F-MAMMA1001280	0. 1
	MAMMA1001298	F-MAMMA1001298	0.07
	MAMMA1001322	F-MAMMA1001322	0.18
	MAMMA1001324	F-MAMMA1001324	0. 25
25	MAMMA1001330	F-MAMMA1001330	0. 13
	MAMMA1001341	F-MAMMA1001341	0.16
	MAMMA1001383	F-MAMMA1001383	0.05
	MAMMA1001397	F-MAMMA1001397	0. 22
30	MAMMA1001408	F-MAMMA1001408	0.06
	MAMMA1001420	F-MAMMA1001420	0.1
	MAMMA1001435	F-MAMMA1001435	0.07
	MAMMA1001442	F-MAMMA1001442	0.13
35	MAMMA1001446	F-MAMMA1001446	0.08
	MAMMA1001452	F-MAMMA1001452	0.1
	MAMMA1001501	F-MAMMA1001501	0.12
	MAMMA1001502	F-MAMMA1001502	0.09
40	MAMMA1001547	F-MAMMA1001547	0. 22
	MAMMA1001575	F-MAMMA1001575	0.08
	MAMMA1001590	F-MAMMA1001590	0.14
	MAMMA1001606	F-MAMMA1001606	0.13
	MAMMA1001663	F-MAMMA1001663	0.11
45	MAMMA1001670	F-MAMMA1001670	0. 2
	MAMMA1001671	F-MAMMA1001671	0.13
	MAMMA1001679	F-MAMMA1001679	0.13
	MAMMA1001683	F-MAMMA1001683	0.16
50	MAMMA1001686	F-MAMMA1001686	0.09
	MAMMA1001711	F-MAMMA1001711	0.11
	MAMMA1001715	F-MAMMA1001715	0.11
	MAMMA1001740	F-MAMMA1001740	0. 1
55	MAMMA1001745	F-MAMMA1001745	0.09

	MAMMA1001760	F-MAMMA1001760	0. 16
	MAMMA1001769	F-MAMMA1001769	0.09
	MAMMA1001788	F-MAMMA1001788	0.03
5	MAMMA1001806	F-MAMMA1001806	
	MAMMA1001815	F-MAMMA1001800	0.1
	MAMMA1001818		
		F-MAMMA1001818	0.1
10	MAMMA1001820	F-MAMMA1001820	0. 27
10	MAMMA1001836	F-MAMMA1001836	0.05
	MAMMA1001854	F-MAMMA1001854	0. 17
	MAMMA1001878	F-MAMMA1001878	0. 13
	MAMMA1001880	F-MAMMA1001880	0. 15
15	MAMMA1001890	F-MAMMA1001890	0. 05
	MAMMA1001907	F-MAMMA1001907	0. 1
	MAMMA1001908	F-MAMMA1001908,	
	MAMMA1001931	F-MAMMA1001931	0.14
20	MAMMA1001963	F-MAMMA1001963	0.08
	MAMMA1001992	F-MAMMA1001992	0.12
	MAMMA1002032	F-MAMMA1002032	0.08
	MAMMA1002056	F-MAMMA1002056	0.09
25	MAMMA1002058	F-MAMMA1002058	0.16
	MAMMA1002078	F-MAMMA1002078	0. 28
	MAMMA1002082	F-MAMMA1002082	0.08
	MAMMA1002084	F-MAMMA1002084	0. 24
•	MAMMA1002093	F-MAMMA1002093	0. 12
30	MAMMA1002108	F-MAMMA1002108	0.05
	MAMMA1002118	F-MAMMA1002118	0.1
	MAMMA1002125	F-MAMMA1002125	0.09
	MAMMA1002132	F-MAMMA1002132	0.11
35	MAMMA1002140	F-MAMMA1002140	0.11
	MAMMA1002145	F-MAMMA1002145	0.11
	MAMMA1002155	F-MAMMA1002155	0.1
	MAMMA1002158	F-MAMMA1002158	0.11
40	MAMMA1002215	F-MAMMA1002215	0.05
	MAMMA1002230	F-MAMMA1002230	0.07
	MAMMA1002250	F-MAMMA1002250	0. 2
	MAMMA1002267	F-MAMMA1002267	0.08
45	MAMMA1002282	F-MAMMA1002282	0.05
	MAMMA1002293	F-MAMMA1002293	0.09
	MAMMA1002298	F-MAMMA1002298	0.09
	MAMMA1002299	F-MAMMA1002299	0. 16
50	MAMMA1002310	F-MAMMA1002310	0. 16
50	MAMMA1002311	F-MAMMA1002311	0. 13
	MAMMA1002322	F-MAMMA1002322	0.07
	MAMMA1002332	F-MAMMA1002332	0.08
	MAMMA1002339	F-MAMMA1002339	0.09
55	MAMMA1002347	F-MAMMA1002347	0.06

	MAMMA1002352	F-MAMMA1002352	0.08
	MAMMA1002359	F-MAMMA1002359	0.13
	MAMMA1002360	F-MAMMA1002360	0.17
5	MAMMA1002361	F-MAMMA1002361	0.13
	MAMMA1002392	F-MAMMA1002392	0.16
	MAMMA1002411	F-MAMMA1002411	0. 28
	MAMMA1002417	F-MAMMA1002417	0.19
10	MAMMA1002428	F-MAMMA1002428	0.07
	MAMMA1002446	F-MAMMA1002446	0. 17
	MAMMA1002475	F-MAMMA1002475	0.17
	MAMMA1002480	F-MAMMA1002480	0.07
15	MAMMA1002494	F-MAMMA1002494	0.17
	MAMMA1002498	F-MAMMA1002498	0. 28
	MAMMA1002545	F-MAMMA1002545,	0.12
	MAMMA1002556	F-MAMMA1002556	0.06
20	MAMMA1002566	F-MAMMA1002566	0.13
	MAMMA1002571	F-MAMMA1002571	0. 29
	MAMMA1002573	F-MAMMA1002573	0. 29
	MAMMA1002597	F-MAMMA1002597	0.1
25	MAMMA1002603	F-MAMMA1002603	0.1
	MAMMA1002612	F-MAMMA1002612	0.14
	MAMMA1002618	F-MAMMA1002618	0. 11
	MAMMA1002622	F-MAMMA1002622	0. 24
30	MAMMA1002623	,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	0. 1
30	MAMMA1002625	F-MAMMA1002625	0. 11
	MAMMA1002629	F-MAMMA1002629	0.05
	MAMMA1002636	F-MAMMA1002636	0. 12
	MAMMA1002646	F-MAMMA1002646	0.05
35	MAMMA1002662	F-MAMMA1002662	0.09
	MAMMA1002698 MAMMA1002701	F-MAMMA1002698 F-MAMMA1002701	0.08
	MAMMA1002701		0. 11
	MAMMA1002708	F-MAMMA1002708 F-MAMMA1002721	0.09
40	MAMMA1002727	F-MAMMA1002727	0. 06 0. 15
	MAMMA1002727	F-MAMMA1002728	0. 13
	MAMMA1002744	F-MAMMA1002744	0. 12
	MAMMA1002746	F-MAMMA1002746	0.09
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	MAMMA1002754	F-MAMMA1002754	0. 07
	MAMMA1002758	F-MAMMA1002758	0. 29
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	MAMMA1002780	F-MAMMA1002780	0.12
	MAMMA1002820	F-MAMMA1002820	0. 3
	MAMMA1002830	F-MAMMA1002830	0.16
55	MAMMA1002833	F-MAMMA1002833	0.14

	MAMMA1002835	F-MAMMA1002835	0. 13
	MAMMA1002838	F-MAMMA1002838	0.09
	MAMMA1002844	F-MAMMA1002844	0. 27
5	MAMMA1002858	F-MAMMA1002858	
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	MAMMA1002880	F-MAMMA1002880	0.11
	MAMMA1002887	F-MAMMA1002887	0.09
10	MAMMA1002892	F-MAMMA1002892	0. 17
	MAMMA1002908	F-MAMMA1002908	0.11
	MAMMA1002909	F-MAMMA1002909	0. 22
	MAMMA1002930	F-MAMMA1002930	0. 2
15	MAMMA1002941	F-MAMMA1002941	0. 23
	MAMMA1002970	F-MAMMA1002970	0.06
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	MAMMA1002973	F-MAMMA1002973	0.1
20	MAMMA1002982	F-MAMMA1002982	0. 1
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	MAMMA1003007	F-MAMMA1003007	0.09
	MAMMA1003019	F-MAMMA1003019	0.12
25	MAMMA1003026	F-MAMMA1003026	0. 26
	MAMMA1003031	F-MAMMA1003031	0.18
	MAMMA1003039	F-MAMMA1003039	0.09
	MAMMA1003040	F-MAMMA1003040	0.07
30	MAMMA1003055	F-MAMMA1003055	0.08
	MAMMA1003089	F-MAMMA1003089	0.14
	MAMMA1003140	F-MAMMA1003140	0.19
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35	NT2RM4000046	F-NT2RM4000046	0.1
	NT2RM4000085	F-NT2RM4000085	0. 23
	NT2RM4000086	F-NT2RM4000086	0. 28
	NT2RM4000200	F-NT2RM4000200	0.16
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	NT2RM4000265	F-NT2RM4000265	0.18
	NT2RM4000327	F-NT2RM4000327	0.06
	NT2RM4000366	F-NT2RM4000366	0. 28
45	NT2RM4000368	F-NT2RM4000368	0.11
•	NT2RM4000414	F-NT2RM4000414	0.19
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	NT2RM4001016	F-NT2RM4001016	0. 2
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	NT2RM4001605	F-NT2RM4001605	0. 22
	NT2RM4001754	F-NT2RM4001754	0. 26
	NT2RM4001776	F-NT2RM4001776	0. 18
10	NT2RM4001810	F-NT2RM4001810	0. 22
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	NT2RM4001858	F-NT2RM4001858	0.14
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	NT2RM4001984	F-NT2RM4001984	0.09
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	NT2RM4002278	F-NT2RM4002278	0.07
20	NT2RM4002281	F-NT2RM4002281	0.07
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	NT2RM4002383	F-NT2RM4002383	0.08
	NT2RM4002390	F-NT2RM4002390	0.18
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	NT2RP2000327	F-NT2RP2000327	0. 14
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	NT2RP2001449	F-NT2RP2001449	0.15
20	NT2RP2001467	F-NT2RP2001467	0. 21
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	NT2RP2001526	F-NT2RP2001526	0.1
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	NT2RP2001678	F-NT2RP2001678	0.17
	NT2RP2001699	F-NT2RP2001699	0.05
	NT2RP2001720	F-NT2RP2001720	0.08
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	NT2RP2001813	F-NT2RP2001813	0. 19
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	NT2RP2002032	F-NT2RP2002032	0. 23
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	NT2RP2002124	F-NT2RP2002124	0.13
	NT2RP2002137	F-NT2RP2002137	0. 07
	NT2RP2002172	F-NT2RP2002172	0. 21
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	NT2RP2002256	F-NT2RP2002256	0.17
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55	NT2RP2002394	F-NT2RP2002394	0.16

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	NT2RP2002457	F-NT2RP2002457	0. 07
	NT2RP2002437	F-NT2RP2002475	0. 11
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	NT2RP2002504	F-NT2RP2002504	• • •
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	NT2RP2002591	F-NT2RP2002591	0. 15
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	NT2RP2002727	F-NT2RP2002727	0. 26
	NT2RP2002736	F-NT2RP2002736	0. 12
	NT2RP2002740	F-NT2RP2002740	0.16
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	NT2RP2002857	F-NT2RP2002857	0. 21
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25	NT2RP2003073	F-NT2RP2003073	0.08
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	NT2RP2003339	F-NT2RP2003339	0.1
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	NT2RP2003456	F-NT2RP2003456	0.08
	NT2RP2003499	F-NT2RP2003499	0.08
	NT2RP2003517	F-NT2RP2003517	0. 29
40	NT2RP2003522	F-NT2RP2003522	0. 22
70	NT2RP2003559	F-NT2RP2003559	0.12
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	NT2RP2003704	F-NT2RP2003704	0.09
	NT2RP2003706	F-NT2RP2003706	0. 28
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	NT2RP2003770	F-NT2RP2003770	0.09
	NT2RP2003859	F-NT2RP2003859	0.18
	NT2RP2003871	F-NT2RP2003871	0. 23
50	NT2RP2003885	F-NT2RP2003885	0. 24
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	NT2RP2003968	F-NT2RP2003968	0. 2
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	NT2RP2004412	F-NT2RP2004412	0. 24
	NT2RP2004425	F-NT2RP2004425	-
	NT2RP2004490	F-NT2RP2004490	0. 13
20	NT2RP2004512	F-NT2RP2004512	0. 10
	NT2RP2004523	F-NT2RP2004523	0. 15
	NT2RP2004580	F-NT2RP2004580	0. 12
	NT2RP2004587	F-NT2RP2004587	0.08
25	NT2RP2004675	F-NT2RP2004675	0.14
20	NT2RP2004681	F-NT2RP2004681	0. 17
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••	NT2RP2004767	F-NT2RP2004767	0. 21
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	NT2RP2005020	F-NT2RP2005020	0.16
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	NT2RP2005289	F-NT2RP2005289	0. 22
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	NT2RP2005857	F-NT2RP2005857	0. 28
15	NT2RP2005908	F-NT2RP2005908	0.05
	NT2RP2005942	F-NT2RP2005942	0. 26
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20	NT2RP2006103	F-NT2RP2006103	0.07
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	NT2RP2006261	F-NT2RP2006261	0.16
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	NT2RP3000451	F-NT2RP3000451	0.14
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	NT2RP3000561	F-NT2RP3000561	0. 1
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55	NT2RP3000582	F-NT2RP3000582	0.08

			
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	NT2RP3000742	F-NT2RP3000742	0. 22
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	NT2RP3000875	F-NT2RP3000875	0. 16
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	NT2RP3001339	F-NT2RP3001339 F-NT2RP3001340	0. 27
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30	NT2RP3001580	F-NT2RP3001580	0. 23
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	NT2RP3001634	F-NT2RP3001634	0. 15
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35	NT2RP3001782	F-NT2RP3001782	0. 21
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40	NT2RP3002002	F-NT2RP3002002	0.18
	NT2RP3002004	F-NT2RP3002004	0.11
	NT2RP3002056	F-NT2RP3002056	0. 16
	NT2RP3002057	F-NT2RP3002057	0.08
45	NT2RP3002166	F-NT2RP3002166	0. 15
	NT2RP3002173	F-NT2RP3002173	0. 28
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	NT2RP3002810	F-NT2RP3002810	0. 2
	NT2RP3002955	F-NT2RP3002955	0. 17
55	NT2RP3002978	F-NT2RP3002978	0.06

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10	NT2RP3003384	F-NT2RP3003384	0.12
	NT2RP3003403	F-NT2RP3003403	0. 26
	NT2RP3003433	F-NT2RP3003433	0. 22
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7.5	NT2RP3003572	F-NT2RP3003572	0. 29
	NT2RP3003576	F-NT2RP3003576,	
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	NT2RP3003665	F-NT2RP3003665	0. 15
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	NT2RP3003800	F-NT2RP3003800	0. 28
	NT2RP3003819	F-NT2RP3003819	0. 19
25	NT2RP3003828	F-NT2RP3003828	0. 12
	NT2RP3003842	F-NT2RP3003842	0.11
	NT2RP3003989	F-NT2RP3003989	0. 1
	NT2RP3004016	F-NT2RP3004016	0.09
30	NT2RP3004070	F-NT2RP3004070	0. 11
	NT2RP3004095	F-NT2RP3004095	0. 1
	NT2RP3004110	F-NT2RP3004110	0.07
	NT2RP3004145	F-NT2RP3004145	0.15
35	NT2RP3004148	F-NT2RP3004148	0.14
	NT2RP3004215	F-NT2RP3004215	0.16
	NT2RP3004334	F-NT2RP3004334	0. 1
	NT2RP3004349	F-NT2RP3004349	0.09
40	NT2RP3004399	F-NT2RP3004399	0.16
70	NT2RP3004470	F-NT2RP3004470	0. 23
	NT2RP3004475	F-NT2RP3004475	0. 26
	NT2RP3004503	F-NT2RP3004503	0.12
_	NT2RP3004527	F-NT2RP3004527	0.06
45	NT2RP4000023	F-NT2RP4000023	0.13
	NT2RP4000035	F-NT2RP4000035	0. 17
	NT2RP4000049	F-NT2RP4000049	0. 1
	NT2RP4000078	F-NT2RP4000078	0. 28
50	NT2RP4000102	F-NT2RP4000102	0. 15
	NT2RP4000167	F-NT2RP4000167	0.19
	NT2RP4000214	F-NT2RP4000214	0.12
	NT2RP4000218	F-NT2RP4000218	0. 17
55	NT2RP4000321	F-NT2RP4000321	0.18

	NT2RP4000424	F-NT2RP4000424	0. 22
	NT2RP4000515	F-NT2RP4000515	0. 13
5	NT2RP4000517	F-NT2RP4000517	0. 1
	NT2RP4000519	F-NT2RP4000519	0. 26
	NT2RP4000915	F-NT2RP4000915	0. 19
	NT2RP4000996	F-NT2RP4000996	0. 05
40	NT2RP4001407	F-NT2RP4001407	0. 2
10	NT2RP4001889	F-NT2RP4001889	0. 09
	NT2RP4002075	F-NT2RP4002075	0. 19
	NT2RP4002905	F-NT2RP4002905	0.15
	OVARC1000017	F-0VARC1000017	0. 12
15	OVARC1000058	F-0VARC1000058	0. 2
	OVARC1000068	F-0VARC1000068	0. 14
	OVARC1000071	F-0VARC1000071,	0.15
	OVARC1000085	F-0VARC1000085	0. 16
20	OVARC1000092	F-0VARC1000092	0. 26
	OVARC1000133	F-0VARC1000133	0.14
	0VARC1000145	F-0VARC1000145	0. 15
	OVARC1000191	F-0VARC1000191	0.07
25	OVARC1000198	F-0VARC1000198	0. 23
	OVARC1000240	F-0VARC1000240	0.11
	0VARC1000302	F-0VARC1000302	0.08
	0VARC1000414	F-0VARC1000414	0. 12
30	OVARC1000427	F-0VARC1000427	0. 1
	0VARC1000431	F-0VARC1000431	0.06
	OVARC1000486	F-0VARC1000486	0. 08
	OVARC1000496	F-0VARC1000496	0. 13
25	OVARC1000526	F-0VARC1000526	0.09
35	OVARC1000533	F-0VARC1000533	0.11
	OVARC1000543	F-0VARC1000543	0. 2
	OVARC1000573	F-0VARC1000573	0.09
	OVARC1000578	F-0VARC1000578	0.13
40	0VARC1000678	F-0VARC1000678	0.17
	OVARC1000679	F-0VARC1000679	0. 23
	0VARC1000700	F-0VARC1000700	0. 28
	OVARC1000769	F-0VARC1000769	0.05
45	OVARC1000802	F-0VARC1000802	0.06
	0VARC1000891	F-0VARC1000891	0.08
	OVARC1000897 OVARC1000936	F-0VARC1000897 F-0VARC1000936	0.11
	0VARC1000936	F-0VARC1000936	0.14
50	OVARC1000937	F-0VARC1000937	0. 29
	0VARC1000948	F-0VARC1000948	0. 14
	OVARC1000900	F-0VARC1000980	0. 07
	0VARC1000971	F-0VARC1000971	0. 11
55	0VARC1001000	F-0VARC1001000	0. 06
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	0VARC1001051	F-0VARC1001051	0.16
	OVARC1001062	F-0VARC1001062	0. 09
	OVARC1001072	F-0VARC1001072	0.11
5	0VARC1001085	F-0VARC1001085	0. 12
	0VARC1001117	F-0VARC1001117	0. 21
	OVARC1001118	F-0VARC1001118	0. 08
	OVARC1001129	F-0VARC1001129	0. 28
10	OVARC1001169	F-0VARC1001169	0. 12
	OVARC1001240	F-0VARC1001240	0.09
	OVARC1001261	F-0VARC1001261	0. 1
	OVARC1001268	F-0VARC1001268	0. 05
15	OVARC1001282	F-0VARC1001282	0.09
	OVARC1001330	F-0VARC1001330	0. 08
	OVARC1001339	F-0VARC1001339,	0. 15
	OVARC1001342	F-0VARC1001342	0. 16
20	OVARC1001357	F-0VARC1001357	0.06
20	OVARC1001442	F-0VARC1001442	0. 15
	OVARC1001480	F-0VARC1001480	0. 23
	0VARC1001542	F-0VARC1001542	0. 13
	OVARC1001547	F-0VARC1001547	0.12
25	OVARC1001600	F-0VARC1001600	0.13
	OVARC1001615	F-0VARC1001615	0.11
	OVARC1001668	F-0VARC1001668	0. 05
	OVARC1001726	F-0VARC1001726	0.18
30	OVARC1001745	F-0VARC1001745	0. 22
	OVARC1001795	F-0VARC1001795	0. 28
	OVARC1001805	F-0VARC1001805	0. 16
	OVARC1001812	F-0VARC1001812	0. 05
35	OVARC1001813	F-0VARC1001813	0.11
	OVARC1001846	F-0VARC1001846	0. 13
	OVARC1001883	F-0VARC1001883	0. 05
	OVARC1001900	F-0VARC1001900	0. 28
40	OVARC1001901	F-0VARC1001901	0.09
	OVARC1001911	F-0VARC1001911	0. 06
	OVARC1001928	F-0VARC1001928	0. 1
	OVARC1001989	F-0VARC1001989	
45	0VARC1002044	F-0VARC1002044	0.06
**	OVARC1002066	F-0VARC1002066	0. 08
	OVARC1002143	F-0VARC1002143	0. 27
	PLACE1000031	F-PLACE1000031	0.09
	PLACE1000040	F-PLACE1000040	0. 15
50	PLACE1000078	F-PLACE1000078	0. 12
	PLACE1000094	F-PLACE1000094	0.16
	PLACE1000214	F-PLACE1000214	0. 15
	PLACE1000292	F-PLACE1000292	0. 05
55	PLACE1000332	F-PLACE1000332	0. 1

	PLACE1000421	F-PLACE1000421	0.14
	PLACE1000424	F-PLACE1000424	0.08
	PLACE1000481	F-PLACE1000481	0. 12
5	PLACE1000540	F-PLACE1000540	0. 12
	PLACE1000583	F-PLACE1000583	0.11
	PLACE1000599	F-PLACE1000585	0. 1
	PLACE1000599		0. 22
10		F-PLACE1000749	
	PLACE1000841	F-PLACE1000841	0.09
	PLACE1000931	F-PLACE1000931	0.22
	PLACE1000979	F-PLACE1000979	0.16
	PLACE1001007	F-PLACE1001007	0.19
15	PLACE1001015	F-PLACE1001015	0. 25
	PLACE1001076	F-PLACE1001076	0.08
	PLACE1001088	F-PLACE1001088,	
	PLACE1001118	F-PLACE1001118	0. 24
20	PLACE1001136	F-PLACE1001136	0.12
	PLACE1001272	F-PLACE1001272	0.07
	PLACE1001279	F-PLACE1001279	0.08
	PLACE1001323	F-PLACE1001323	0.18
25	PLACE1001377	F-PLACE1001377	0.16
	PLACE1001384	F-PLACE1001384	0.06
	PLACE1001395	F-PLACE1001395	0.08
	PLACE1001414	F-PLACE1001414	0.11
30	PLACE1001440	F-PLACE1001440	0.14
30	PLACE1001468	F-PLACE1001468	0.05
	PLACE1001502	F-PLACE1001502	0.06
	PLACE1001517	F-PLACE1001517	0.09
	PLACE1001534	F-PLACE1001534	0.08
35	PLACE1001603	F-PLACE1001603	0.08
	PLACE1001611	F-PLACE1001611	0. 25
	PLACE1001640 PLACE1001672	F-PLACE1001640	0.09
	PLACE1001691	F-PLACE1001672 F-PLACE1001691	0. 11 0. 11
40	PLACE1001091	F-PLACE1001705	0. 11
	PLACE1001705	F-PLACE1001705	
	PLACE1001770	F-PLACE1001718	0. 15 0. 18
	PLACE1001745	F-PLACE1001720	0. 18
45	PLACE1001746	F-PLACE1001746	0. 09
	PLACE1001748	F-PLACE1001748	0. 12
	PLACE1001756	F-PLACE1001748	0. 17
	PLACE1001799	F-PLACE1001799	0.17
50	PLACE1001821	F-PLACE1001799	0. 14
	PLACE1001845	F-PLACE1001845	0.03
	PLACE1001897	F-PLACE1001897	0. 08
	PLACE1002052	F-PLACE1002052	0. 13
	PLACE1002066	F-PLACE1002066	0. 13
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	PLACE1002150	F-PLACE1002150	0. 08
	PLACE1002157	F-PLACE1002157	0. 15
	PLACE1002205	F-PLACE1002205	0. 13
5	PLACE1002227	F-PLACE1002227	0. 12
	PLACE1002256	F-PLACE1002256	0.11
	PLACE1002259	F-PLACE1002259	0. 09
	PLACE1002319	F-PLACE1002319	0. 27
10	PLACE1002399	F-PLACE1002399	0. 08
	PLACE1002477	F-PLACE1002477	0. 17
	PLACE1002493	F-PLACE1002493	0. 09
	PLACE1002500	F-PLACE1002500	0. 25
15	PLACE1002514	F-PLACE1002514	0. 06
	PLACE1002537	F-PLACE1002537	0. 17
	PLACE1002578	F-PLACE1002578,	0. 11
	PLACE1002583	F-PLACE1002583	0. 25
20	PLACE1002591	F-PLACE1002591	0. 28
20	PLACE1002598	F-PLACE1002598	0. 29
	PLACE1002604	F-PLACE1002604	0. 15
	PLACE1002768	F-PLACE1002768	0. 11
	PLACE1002772	F-PLACE1002772	0. 15
25	PLACE1002782	F-PLACE1002782	0. 16
	PLACE1002853	F-PLACE1002853	0. 16
	PLACE1002881	F-PLACE1002881	0.09
	PLACE1002962	F-PLACE1002962	0.13
30	PLACE1002968	F-PLACE1002968	0. 23
	PLACE1003153	F-PLACE1003153	0. 07
	PLACE1003205	F-PLACE1003205	0. 2
	PLACE1003238	F-PLACE1003238	0. 22
35	PLACE1003258	F-PLACE1003258	0.17
	PLACE1003343	F-PLACE1003343	0. 07
	PLACE1003361	F-PLACE1003361	0. 19
	PLACE1003373	F-PLACE1003373	0.08
40	PLACE1003375	F-PLACE1003375	0. 26
	PLACE1003401	F-PLACE1003401	0.11
	PLACE1003420	F-PLACE1003420	0. 22
	PLACE1003478	F-PLACE1003478	0.14
45	PLACE1003516	F-PLACE1003516	0.17
45	PLACE1003519	F-PLACE1003519	0.14
	PLACE1003521	F-PLACE1003521	0. 17
	PLACE1003528	F-PLACE1003528	0. 16
	PLACE1003566	F-PLACE1003566	0. 16
50	PLACE1003575	F-PLACE1003575	0.06
	PLACE1003583	F-PLACE1003583	0.08
	PLACE1003584	F-PLACE1003584	0. 1
	PLACE1003593	F-PLACE1003593	0. 15
55	PLACE1003618	F-PLACE1003618	0. 25

	PLACE1003638	F-PLACE1003638	0.06
	PLACE1003760	F-PLACE1003760	0.12
	PLACE1003768	F-PLACE1003768	0. 13
5	PLACE1003833	F-PLACE1003833	0.06
	PLACE1003850	F-PLACE1003850	0. 07
	PLACE1003858	F-PLACE1003858	0. 07
	PLACE1003864	F-PLACE1003864	0. 12
10	PLACE1003900	F-PLACE1003900	0. 11
	PLACE1003932	F-PLACE1003932	0.05
	PLACE1004118	F-PLACE1004118	0. 3
	PLACE1004242	F-PLACE1004242	0. 05
15	PLACE1004256	F-PLACE1004256	0. 24
	PLACE1004257	F-PLACE1004257	0.06
	PLACE1004274	F-PLACE1004274,	0. 25
	PLACE1004284	F-PLACE1004284	0.13
20	PLACE1004336	F-PLACE1004336	0. 27
	PLACE1004384	F-PLACE1004384	0.17
	PLACE1004425	F-PLACE1004425	0.12
	PLACE1004467	F-PLACE1004467	0. 2
	PLACE1004471	F-PLACE1004471	0.12
25	PLACE1004491	F-PLACE1004491	0.08
	PLACE1004506	F-PLACE1004506	
	PLACE1004518	F-PLACE1004518	0.14
	PLACE1004658	F-PLACE1004658	0.05
30	PLACE1004681	F-PLACE1004681	0.13
	PLACE1004693	F-PLACE1004693	0.11
	PLACE1004716	F-PLACE1004716	0.12
	PLACE1004773	F-PLACE1004773	0.17
35	PLACE1004793	F-PLACE1004793	0. 21
	PLACE1004813	F-PLACE1004813	0.11
	PLACE1004815	F-PLACE1004815	0.14
	PLACE1004827	F-PLACE1004827	0.15
40	PLACE1004836	F-PLACE1004836	0. 25
	PLACE1004838	F-PLACE1004838	0.16
	PLACE1004840	F-PLACE1004840	0.06
	PLACE1004900	F-PLACE1004900	0.08
45	PLACE1004913	F-PLACE1004913	0. 12
40	PLACE1004934	F-PLACE1004934	0.17
	PLACE1004972	F-PLACE1004972	0.07
	PLACE1004979	F-PLACE1004979	0. 24
	PLACE1004985	F-PLACE1004985	0.05
50	PLACE1005052	F-PLACE1005052	0.17
	PLACE1005077	F-PLACE1005077	0.1
	PLACE1005085 PLACE1005086	F-PLACE1005085 F-PLACE1005086	0.13
	PLACE 1005086	F-PLACE1005086	0.13
55	1 EVOT 1009 108	F-FLAGE 1003 108	0. 28

	PLACE1005111	F-PLACE1005111	0. 15
	PLACE1005128	F-PLACE1005128	0.12
	PLACE1005146	F-PLACE1005146	0. 07
5	PLACE1005162	F-PLACE1005162	0. 13
	PLACE1005176	F-PLACE1005176	0.16
	PLACE1005409	F-PLACE1005409	0. 09
	PLACE1005453	F-PLACE1005453	0.1
10	PLACE1005467	F-PLACE1005467	0. 28
	PLACE1005471	F-PLACE1005471	0.06
	PLACE1005477	F-PLACE1005477	0.12
	PLACE1005502	F-PLACE1005502	0. 1
15	PLACE1005526	F-PLACE1005526	
	PLACE1005528	F-PLACE1005528	0. 07
	PLACE1005574	F-PLACE1005574,	0. 06
	PLACE1005584	F-PLACE1005584	0. 24
22	PLACE1005611	F-PLACE1005611	0. 18
20	PLACE1005639	F-PLACE1005639	0. 09
	PLACE1005666	F-PLACE1005666	0. 07
·	PLACE1005730	F-PLACE1005730	0. 2
	PLACE1005802	F-PLACE1005802	0. 1
25	PLACE1005845	F-PLACE1005845	0. 09
	PLACE1005850	F-PLACE1005850	0. 22
	PLACE1005884	F-PLACE1005884	0. 08
	PLACE1005898	F-PLACE1005898	0.12
30	PLACE1005932	F-PLACE1005932	0. 13
	PLACE1005968	F-PLACE1005968	0. 07
	PLACE1006002	F-PLACE1006002	0. 08
	PLACE1006003	F-PLACE1006003	0. 16
35	PLACE1006017	F-PLACE1006017	0. 13
	PLACE1006076	F-PLACE1006076	0. 11
	PLACE1006129	F-PLACE1006129	0. 25
	PLACE1006143	F-PLACE1006143	0. 08
40	PLACE1006164	F-PLACE1006164	0. 05
	PLACE1006187	F-PLACE1006187	0. 2
	PLACE1006205	F-PLACE1006205	0. 08
		F-PLACE1006223	0. 05
45		F-PLACE1006262	0. 08
	PLACE1006288	F-PLACE1006288	0.06
	PLACE1006318	F-PLACE1006318	0. 12
	PLACE1006360	F-PLACE1006360	0. 07
50	PLACE1006368	F-PLACE1006368	0. 27
50	PLACE1006371	F-PLACE1006371	0. 14
	PLACE1006382	F-PLACE1006382	0.09
	PLACE1006470 PLACE1006506	F-PLACE1006470 F-PLACE1006506	0. 14 0. 24
	PLACE1006506	F-PLACE1006506	0. 24
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	PLACE1006534	F-PLACE1006534	0.06
	PLACE1006540	F-PLACE1006540	0. 05
	PLACE1006617	F-PLACE1006617	0. 05
5	PLACE1006629	F-PLACE1006629	0. 2
	PLACE1006640	F-PLACE1006640	0. 05
	PLACE1006754	F-PLACE1006754	0. 08
	PLACE1006760	F-PLACE1006760	0. 05
10	PLACE1006779	F-PLACE1006779	0. 17
,-	PLACE1006792	F-PLACE1006792	0. 15
	PLACE1006795	F-PLACE1006795	0. 1
	PLACE1006800	F-PLACE1006800	0. 11
	PLACE1006805	F-PLACE1006805	0. 11
15	PLACE1006815	F-PLACE1006815	0. 12
	PLACE1006819	F-PLACE1006860,	
	PLACE1006867	F-PLACE1006867	
	PLACE1006807	F-PLACE1006878	0. 11 0. 23
20	PLACE1006976		
	PLACE1008904 PLACE1007045	F-PLACE1006904 F-PLACE1007045	0. 13
	PLACE1007045 PLACE1007097	F-PLACE1007097	0. 08
	PLACE1007097	F-PLACE1007097	0. 13
25	PLACE1007111	F-PLACE1007111	0. 08
	PLACE1007132	F-PLACE1007140	0.07
			0. 16
	PLACE1007276	F-PLACE1007276	
••	PLACE1007286 PLACE1007478	F-PLACE1007286	0. 05 0. 19
30	PLACE1007478 PLACE1007525	F-PLACE1007478 F-PLACE1007525	0. 19
	PLACE1007525	F-PLACE1007557	0. 11
	PLACE1007677		
	PLACE1007677 PLACE1007737	F-PLACE1007677 F-PLACE1007737	0. 08 0. 07
35	PLACE1007737	F-PLACE1007737	0.07
	PLACE1007/43	F-PLACE1007743	0. 09
	PLACE1007829	F-PLACE1007829	0. 09
	PLACE1007852	F-PLACE1007852	0. 09
40	PLACE1007866	F-PLACE1007866	0. 14
	PLACE1007877	F-PLACE1007877	0. 07
	PLACE1007877	F-PLACE1008045	0. 29
	PLACE1008045	F-PLACE1008080	
45	PLACE1008080	F-PLACE1008080	0.05
	PLACE1008111	F-PLACE1008111	0.11
	PLACE1008181	F-PLACE1008181	0.08
	PLACE1008201	F-PLACE1008201	0. 12
	PLACE1008231		0.08
50		F-PLACE1008244	0.06
	PLACE1008330	F-PLACE1008330	0.13
	PLACE1008331	F-PLACE1008331	0.15
,	PLACE1008369	F-PLACE1008369	0.06
55	PLACE1008392	F-PLACE1008392	0. 09

		_	
	PLACE1008405	F-PLACE1008405	0.07
	PLACE1008424	F-PLACE1008424	0.06
	PLACE1008426	F-PLACE1008426	0. 17
5	PLACE1008437	F-PLACE1008437	0. 2
	PLACE1008455	F-PLACE1008455	0.07
	PLACE1008584	F-PLACE1008584	0. 29
	PLACE1008621	F-PLACE1008621	0.07
10	PLACE1008625	F-PLACE1008625	0.12
	PLACE1008630	F-PLACE1008630	0.12
	PLACE1008643	F-PLACE1008643	0.11
	PLACE1008715	F-PLACE1008715	0. 2
15	PLACE1008748	F-PLACE1008748	0.15
	PLACE1008757	F-PLACE1008757	0.13
	PLACE1008798	F-PLACE1008798,	0.14
	PLACE1008807	F-PLACE1008807	0.12
20	PLACE1008851	F-PLACE1008851	0. 22
	PLACE1008920	F-PLACE1008920	0.1
	PLACE1008941	F-PLACE1008941	0.14
	PLACE1008947	F-PLACE1008947	0.05
25	PLACE1009020	F-PLACE1009020	0.14
25	PLACE1009039	F-PLACE1009039	0.16
	PLACE1009048	F-PLACE1009048	0.09
	PLACE1009050	F-PLACE1009050	0.09
	PLACE1009150	F-PLACE1009150	0.12
30	PLACE1009155	F-PLACE1009155	0.08
	PLACE1009172	F-PLACE1009172	0.07
	PLACE1009183	F-PLACE1009183	0.07
	PLACE1009200 PLACE1009246	F-PLACE1009200 F-PLACE1009246	0.08
35	PLACE1009248 PLACE1009308	F-PLACE1009248	0. 08
	PLACE1009398	F-PLACE1009398	0. 26
	PLACE1009410	F-PLACE1009410	0. 15
	PLACE1009477		0.18
40	PLACE1009493	F-PLACE1009493	0. 23
	PLACE1009539		
	PLACE1009595	F-PLACE1009595	0.11
	PLACE1009613	F-PLACE1009613	0.08
45	PLACE1009621	F-PLACE1009621	0.19
	PLACE1009637	F-PLACE1009637	0. 27
	PLACE1009639	F-PLACE1009639	0. 27
	PLACE1009879	F-PLACE1009879	0. 07
50	PLACE1009888	F-PLACE1009888	0.06
	PLACE1009924	F-PLACE1009924	0.09
	PLACE1009925	F-PLACE1009925	0. 21
	PLACE1009935	F-PLACE1009935	0. 22
55	PLACE1009947	F-PLACE1009947	0.19

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	PLACE1010069	F-PLACE1010069	0. 22
	PLACE1010083	F-PLACE1010083	0. 26
	PLACE1010089	F-PLACE1010089	0. 13
5	PLACE1010181	F-PLACE1010181	0, 07
	PLACE1010231	F-PLACE1010231	0. 17
	PLACE1010270	F-PLACE1010270	0.08
	PLACE1010341	F-PLACE1010341	0.1
10	PLACE1010562	F-PLACE1010562	0.14
	PLACE1010579	F-PLACE1010579	0.15
	PLACE1010624	F-PLACE1010624	0.13
	PLACE1010628	F-PLACE1010628	0.16
15	PLACE1010631	F-PLACE1010631	0. 23
	PLACE1010662	F-PLACE1010662	0. 24
	PLACE1010702	F-PLACE1010702,	0. 21
	PLACE1010739	F-PLACE1010739	0. 13
	PLACE1010802	F-PLACE1010802	0. 17
20	PLACE1010891	F-PLACE1010891	0.09
	PLACE1010896	F-PLACE1010896	0. 22
	PLACE1010916	F-PLACE1010916	0.15
	PLACE1010947	F-PLACE1010947	0.08
25	PLACE1011032	F-PLACE1011032	0.17
	PLACE1011057	F-PLACE1011057	0. 1
	PLACE1011109	F-PLACE1011109	0.08
	PLACE1011133	F-PLACE1011133	0.08
30	PLACE1011143	F-PLACE1011143	0. 23
	PLACE1011165	F-PLACE1011165	0. 05
	PLACE1011185	F-PLACE1011185	0. 17
	PLACE1011203	F-PLACE1011203	0. 08
35	PLACE1011296	F-PLACE1011296	0. 13
33	PLACE1011340	F-PLACE1011340	0. 05
	PLACE1011375	F-PLACE1011375	0. 07
	PLACE1011419	F-PLACE1011419	0. 15
	PLACE1011452	F-PLACE1011452	0. 13
40	PLACE1011465	F-PLACE1011465	0.14
	PLACE1011492	F-PLACE1011492	0. 26
	PLACE1011503	F-PLACE1011503	0.14
	PLACE1011520	F-PLACE1011520	0. 12
45	PLACE1011567	F-PLACE1011567	0. 13
	PLACE1011576	F-PLACE1011576	0. 29
	PLACE1011641	F-PLACE1011641	0. 07
	PLACE1011643	F-PLACE1011643	0. 27
50	PLACE1011649	F-PLACE1011649	0. 17
	PLACE1011650	F-PLACE1011650	0. 13
	PLACE1011719	F-PLACE1011719	0. 15
	PLACE1011729	F-PLACE1011729	0. 23
55	PLACE1011749	F-PLACE1011749	0.12
55	* *		

	PLACE1011762	F-PLACE1011762	0.06
	PLACE1011778	F-PLACE1011778	0. 07
	PLACE1011874	F-PLACE1011874	0. 07
5	PLACE1011962	F-PLACE1011962	0.06
	PLACE1011964	F-PLACE1011964	0. 13
	PLACE1011995	F-PLACE1011995	0.08
	PLACE2000003	F-PLACE2000003	0. 09
10	PLACE2000011	F-PLACE2000011	0. 12
	PLACE2000017	F-PLACE2000017	0. 22
	PLACE2000061	F-PLACE2000061	0. 19
	PLACE2000103	F-PLACE2000103	0. 16
45	PLACE2000115	F-PLACE2000115	0. 11
15	PLACE2000132	F-PLACE2000132	0. 24
	PLACE2000136	F-PLACE2000136,	0. 09
	PLACE2000140	F-PLACE2000140	0. 24
	PLACE2000176	F-PLACE2000176	0. 25
20	PLACE2000187	F-PLACE2000187	0. 16
	PLACE2000216	F-PLACE2000216	0. 15
	PLACE2000264	F-PLACE2000264	0.06
	PLACE2000305	F-PLACE2000305	0.05
25	PLACE2000317	F-PLACE2000317	0. 25
	PLACE2000335	F-PLACE2000335	0.09
	PLACE2000342	F-PLACE2000342	0.07
	PLACE2000347	F-PLACE2000347	0.12
30	PLACE2000366	F-PLACE2000366	0.08
	PLACE2000379	F-PLACE2000379	0. 24
	PLACE2000394	F-PLACE2000394	0. 2
	PLACE2000398	F-PLACE2000398	0. 21
35	PLACE2000419	F-PLACE2000419	0.05
	PLACE2000425	F-PLACE2000425	0.16
	PLACE2000435	F-PLACE2000435	0. 05
	PLACE2000450	F-PLACE2000450	0.12
	PLACE2000455	F-PLACE2000455	0. 15
40	PLACE2000465	F-PLACE2000465	0.12
	PLACE2000477	F-PLACE2000477	0.3
	PLACE3000004	F-PLACE3000004	0. 3
	PLACE3000119	F-PLACE3000119	0. 1
45	PLACE3000124	F-PLACE3000124	0.11
	PLACE3000136	F-PLACE3000136	0.13
	PLACE3000158	F-PLACE3000158	0.06
	PLACE3000169	F-PLACE3000169	0.13
50	PLACE3000199	F-PLACE3000199	0. 06
	PLACE3000207	F-PLACE3000207	0.11
	PLACE3000208	F-PLACE3000208	0. 07
	PLACE3000218	F-PLACE3000218	0. 15
55	PLACE3000220	F-PLACE3000220	0. 05

	PLACE3000230	F-PLACE3000230	0. 23
	PLACE3000230	F-PLACE3000271	0. 23
	PLACE3000271	F-PLACE3000276	0.17
5			
•	PLACE3000341	F-PLACE3000341	
	PLACE3000362	F-PLACE3000362	0. 19
	PLACE3000365	F-PLACE3000365	0.09
	PLACE3000373	F-PLACE3000373	0. 22
10	PLACE3000388	F-PLACE3000388	0.12
	PLACE3000399	F-PLACE3000399	0.14
	PLACE3000400	F-PLACE3000400	0. 08
	PLACE3000401	F-PLACE3000401	0. 1
15	PLACE3000402	F-PLACE3000402	0. 1
	PLACE3000406	F-PLACE3000406	0. 17
	PLACE3000475	F-PLACE3000475,	0. 07
	PLACE4000093	F-PLACE4000093	0. 21
20	PLACE4000100	F-PLACE4000100	0.19
	PLACE4000233	F-PLACE4000233	0.07
	PLACE4000247	F-PLACE4000247	0.14
	PLACE4000250	F-PLACE4000250	0.17
	PLACE4000252	F-PLACE4000252	0.09
25	PLACE4000320	F-PLACE4000320	0. 23
	PLACE4000344	F-PLACE4000344	0. 21
	PLACE4000367	F-PLACE4000367	0.18
	PLACE4000379	F-PLACE4000379	0.14
30	PLACE4000401	F-PLACE4000401	0.13
	PLACE4000411	F-PLACE4000411	0. 11
	PLACE4000494	F-PLACE4000494	0.19
	PLACE4000548	F-PLACE4000548	0. 27
35	THYR01000026	F-THYR01000026	0.09
	THYR01000035	F-THYR01000035	0.09
	THYR01000085	F-THYR01000085	0. 13
	THYR01000092	F-THYR01000092	0. 21
40	THYR01000111	F-THYR01000111	0.13
40	THYR01000129	F-THYR01000129	0. 11
	THYR01000132	F-THYR01000132	0. 1
	THYR01000156	F-THYR01000156	0.13
	THYR01000186	F-THYR01000186	0. 12
45	THYR01000187	F-THYR01000187	0. 11
	THYR01000190	F-THYR01000190	0. 16
	THYR01000221	F-THYR01000221	0.08
	THYR01000241	F-THYR01000241	0.08
50	THYR01000279	F-THYR01000279	0.13
	THYR01000438	F-THYR01000438	0.06
	THYR01000452	F-THYR01000452	0. 12
	THYR01000471	F-THYR01000471	0.11
55	THYR01000484	F-THYR01000484	0. 24
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	THYR01000558	F-THYR01000558	0.09
	THYR01000596	F-THYR01000596	0. 14
	THYR01000602	F-THYR01000602	0. 14
5			
	THYR01000625	F-THYR01000625	0.2
	THYR01000641	F-THYR01000641	0.06
	THYR01000658	F-THYR01000658	0. 12
	THYR01000699	F-THYR01000699	0.09
10	THYR01000748	F-THYR01000748	0.3
	THYR01000793	F-THYR01000793	0. 25
	THYR01000796	F-THYR01000796	0.11
	THYR01000805	F-THYR01000805	0. 28
15	THYR01000815	F-THYR01000815	0.14
	THYR01000843	F-THYR01000843	0.13
	THYR01000852	F-THYR01000852,	0.05
	THYR01000865	F-THYR01000865	0.14
20	THYR01000895	F-THYR01000895	0.12
20	THYR01000974	F-THYR01000974	0. 07
	THYR01000975	F-THYR01000975	0.08
	THYR01001003	F-THYR01001003	0.11
	THYR01001031	F-THYR01001031	0.1
25	THYR01001062	F-THYR01001062	0.14
	THYR01001093	F-THYR01001093	0.13
	THYR01001121	F-THYR01001121	0.14
	THYR01001133	F-THYR01001133	0. 24
30	THYR01001177	F-THYR01001177	
	THYR01001262	F-THYR01001262	0. 2
	THYR01001290	F-THYR01001290	0.06
	THYR01001321	F-THYR01001321	0.09
35	THYR01001322	F-THYR01001322	0.13
	THYR01001365	F-THYR01001365	0.08
	THYR01001401	F-THYR01001401	0. 21
	THYR01001411	F-THYR01001411	0.06
	THYR01001426	F-THYR01001426	0.1
40	THYR01001434	F-THYR01001434	0. 15
	THYR01001480	F-THYR01001480	0. 1
	THYR01001534	F-THYR01001534	0. 26
	THYR01001541	F-THYR01001541	0. 22
45	THYR01001559	F-THYR01001559	0.15
	THYR01001570	F-THYR01001570	0. 28
	THYR01001573	F-THYR01001573	0.05
	THYR01001595	F-THYR01001595	0.05
50	THYR01001617	F-THYR01001617	0. 1
	THYR01001673	F-THYR01001673	0.07
	THYR01001706	F-THYR01001706	0. 15
	THYR01001745	F-THYR01001745	0. 12
	THYR01001746	F-THYR01001746	0.1
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	THYR01001772	F-THYR01001772	0.08
	THYR01001895	F-THYR01001895	0.19
5	THYR01001907	F-THYR01001907	0.17
5	Y79AA1000033	F-Y79AA1000033	0.18
	Y79AA1000065	F-Y79AA1000065	0. 22
	Y79AA1000346	F-Y79AA1000346	0. 2
0	Y79AA1000405	F-Y79AA1000405	0. 07
	Y79AA1000410	F-Y79AA1000410	0.08
	Y79AA1000538	F-Y79AA1000538	0.06
	Y79AA1000802	F-Y79AA1000802	0. 28
5	Y79AA1000805	F-Y79AA1000805	0. 2
	Y79AA1000969	F-Y79AA1000969	0. 07
	Y79AA1001061	F-Y79AA1001061	0. 22
	Y79AA1001167	F-Y79AA1001167,	0. 22
)	Y79AA1001384	F-Y79AA1001384	0. 18
	Y79AA1001594	F-Y79AA1001594	0. 22
	Y79AA1001692	F-Y79AA1001692	0. 2
5	Y79AA1001848	F-Y79AA1001848	0. 21
•	Y79AA1001875	F-Y79AA1001875	0. 09
	Y79AA1002103	F-Y79AA1002103	0.11
	Y79AA1002211	F-Y79AA1002211	0.19
0	Y79AA1002220	F-Y79AA1002220	0. 25

Table 11

574 clones belonging to groups (4), (5), (6), and (7), each having the maximal ATGpr1 score of not more than 0.3 and selected based on the maximal ATGpr2 score, and their maximal ATGpr2 scores

10	clone name	name of sequence	maximal ATGpr2 score
15	HEMBA1000180 HEMBA1000251 HEMBA1000280	F-HEMBA1000251 F-HEMBA1000280	0. 37 0. 14 0. 33 0. 36
20	HEMBA1000282 HEMBA1000333 HEMBA1000351 HEMBA1000357 HEMBA1000376	F-HEMBA1000333 F-HEMBA1000351 F-HEMBA1000357	0. 36 0. 59 0. 36 0. 39 0. 53
25	HEMBA1000469 HEMBA1000504 HEMBA1000519	F-HEMBA1000469 F-HEMBA1000504 F-HEMBA1000519	0. 32 0. 34 0. 72
30	HEMBA1000545 HEMBA1000557 HEMBA1000575	F-HEMBA1000557	0. 32 0. 35 0. 34

	HEMBA1000604	F-HEMBA1000604	0. 28
	HEMBA1000622	F-HEMBA1000622	0. 32
_	HEMBA1000673	F-HEMBA1000673	0.14
5	HEMBA1000726	F-HEMBA1000726	0.32
	HEMBA1000749	F-HEMBA1000749	0.35
	HEMBA1000822	F-HEMBA1000822	0. 42
	HEMBA1000876	F-HEMBA1000876	0.44
10	HEMBA1000934	F-HEMBA1000934	0. 41
	HEMBA1000943	F-HEMBA1000943	0.34
	HEMBA1000960	F-HEMBA1000960	0.34
	HEMBA1000972	F-HEMBA1000972	0.44
15	HEMBA1001020	F-HEMBA1001020	0. 31
	HEMBA1001024	F-HEMBA1001024	0. 22
	HEMBA1001026	F-HEMBA1001026,	0. 28
	HEMBA1001051	F-HEMBA1001051	0.36
20	HEMBA1001060	F-HEMBA1001060	0.34
	HEMBA1001071	F-HEMBA1001071	0.57
	HEMBA1001208	F-HEMBA1001208	0. 57
	HEMBA1001226	F-HEMBA1001226	0. 44
25	HEMBA1001319	F-HEMBA1001319	0.58
25	HEMBA1001383	F-HEMBA1001383	0. 35
	HEMBA1001391	F-HEMBA1001391	0. 42
	HEMBA1001411	F-HEMBA1001411	0. 55
	HEMBA1001433	F-HEMBA1001433	0.66
30	HEMBA1001435	F-HEMBA1001435	0.72
	HEMBA1001478	F-HEMBA1001478	0. 36
	HEMBA1001522	F-HEMBA1001522	0.31
	HEMBA1001636	F-HEMBA1001636	0. 32
35	HEMBA1001651	F-HEMBA1001651	0. 54
	HEMBA1001658	F-HEMBA1001658	0. 38
	HEMBA1001709	F-HEMBA1001709	0. 55
	HEMBA1001734	F-HEMBA1001734	0. 13
40	HEMBA1001745	F-HEMBA1001745	0. 33
	HEMBA1001784	F-HEMBA1001784	0. 08
	HEMBA1001791	F-HEMBA1001791	0. 48
	HEMBA1001803	F-HEMBA1001803	0. 21
45	HEMBA1001808	F-HEMBA1001808	0. 23
	HEMBA1001835	F-HEMBA1001835	0. 67
	HEMBA1001844	F-HEMBA1001844	0. 46
	HEMBA1001888	F-HEMBA1001888	0. 72
50	HEMBA1001918	F-HEMBA1001918	0. 25
	HEMBA1001940	F-HEMBA1001940	0. 40
	HEMBA1001962	F-HEMBA1001962	0. 63
	HEMBA1002022	F-HEMBA1002022	0. 32
55	HEMBA1002185	F-HEMBA1002185	0. 33
	HEMBA1002257	F-HEMBA1002257	0. 13

	HEMBA1002348	F-HEMBA1002348	0.39
	HEMBA1002621	F-HEMBA1002621	0. 47
	HEMBA1002645	F-HEMBA1002645	0. 34
5	HEMBA1002661	F-HEMBA1002661	0. 43
	HEMBA1002728	F-HEMBA1002728	0. 45
	HEMBA1002742	F-HEMBA1002742	0.49
	HEMBA1002921	F-HEMBA1002921	0. 79
10	HEMBA1002924	F-HEMBA1002924	0. 45
	HEMBA1002934	F-HEMBA1002934	0.34
	HEMBA1002968	F-HEMBA1002968	0, 39
	HEMBA1003037	F-HEMBA1003037	0.43
15	HEMBA1003064	F-HEMBA1003064	0.19
	HEMBA1003071	F-HEMBA1003071	0.48
•	HEMBA1003166	F-HEMBA1003166,	0.32
	HEMBA1003202	F-HEMBA1003202	0.38
20	HEMBA1003204	F-HEMBA1003204	0, 34
	HEMBA1003220	F-HEMBA1003220	0.39
	HEMBA1003229	F-HEMBA1003229	0.43
	HEMBA1003276	F-HEMBA1003276	0.48
25	HEMBA1003296	F-HEMBA1003296	0.36
	HEMBA1003373	F-HEMBA1003373	0.48
	HEMBA1003531	F-HEMBA1003531	0.46
	HEMBA1003640	F-HEMBA1003640	0.32
20	HEMBA1003714	F-HEMBA1003714	0.16
30	HEMBA1003856	F-HEMBA1003856	0. 47
	HEMBA1003902	F-HEMBA1003902	0.17
	HEMBA1003908	F-HEMBA1003908	0.44
	HEMBA1003926	F-HEMBA1003926	0. 36
35	HEMBA1003942	F-HEMBA1003942	0. 47
	HEMBA1003987	F-HEMBA1003987	0. 39
	HEMBA1004000	F-HEMBA1004000	0. 87
	HEMBA1004024	F-HEMBA1004024	0. 82
40	HEMBA1004164	F-HEMBA1004164	0. 21
	HEMBA1004267	F-HEMBA1004267	0. 46
	HEMBA1004323	F-HEMBA1004323	0. 37
	HEMBA1004405	F-HEMBA1004405	0. 23
45	HEMBA1004433	F-HEMBA1004433	0. 51
	HEMBA1004577	F-HEMBA1004577	0. 51
	HEMBA1004730	F-HEMBA1004730	0. 55
	HEMBA1004778	F-HEMBA1004778	0. 47
50	HEMBA1004803	F-HEMBA1004803	0. 36
	HEMBA1004807	F-HEMBA1004807	0. 48
	HEMBA1004880	F-HEMBA1004880	0. 43
	HEMBA1004900	F-HEMBA1004900	0. 40
55	HEMBA1004909	F-HEMBA1004909	0. 24
	HEMBA1004983	F-HEMBA1004983	0. 35

	HEMBA1005123	F-HEMBA1005123	0. 35
	HEMBA1005232	F~HEMBA1005232	0.09
	HEMBA1005241	F-HEMBA1005241	0. 34
5	HEMBA1005311	F-HEMBA1005311	
	HEMBA1005318	F-HEMBA1005318	0. 59
	HEMBA1005353	F-HEMBA1005353	0. 34
	HEMBA1005374	F-HEMBA1005374	0. 38
10	HEMBA1005447	F-HEMBA1005447	0. 34
	HEMBA1005508	F-HEMBA1005508	0. 14
	HEMBA1005577	F-HEMBA1005577	0. 22
	HEMBA1005588	F-HEMBA1005588	0. 54
15	HEMBA1005593	F-HEMBA1005593	0. 44
	HEMBA1005606	F-HEMBA1005606	0. 38
	HEMBA1005679	F-HEMBA1005679,	
	HEMBA1005894	F-HEMBA1005894	0. 46
20	HEMBA1005911	F-HEMBA1005911	0. 56
	HEMBA1006036	F-HEMBA1006036	0. 87
	HEMBA1006124	F-HEMBA1006124	0. 35
	HEMBA1006253	F-HEMBA1006253	0. 35
25	HEMBA1006259	F-HEMBA1006259	0. 54
	HEMBA1006364	F-HEMBA1006364	0. 55
	HEMBA1006380	F-HEMBA1006380	0. 48
	HEMBA1006426	F-HEMBA1006426	0. 54
30	HEMBA1006461	F-HEMBA1006461	0. 20
	HEMBA1006562	F-HEMBA1006562	0. 72
	HEMBA1006597	F-HEMBA1006597	0. 64
	HEMBA1006639	F-HEMBA1006639	0. 55
25	HEMBA1006653	F-HEMBA1006653	0. 34
35	HEMBA1006695	F-HEMBA1006695	0. 16
	HEMBA1006696	F-HEMBA1006696	0. 37
	HEMBA1006744	F-HEMBA1006744	0. 33
	HEMBA1006824	F-HEMBA1006824	0. 37
40	HEMBA1006949	F-HEMBA1006949	0. 78
	HEMBA1007078	F-HEMBA1007078	0. 44
	HEMBA1007129		
	HEMBA1007147	F-HEMBA1007147	0. 41
45	HEMBA1007206	F-HEMBA1007206	0. 46
	HEMBA1007279	F-HEMBA1007279	0. 45
	HEMBA1007327	F-HEMBA1007327	0. 71
	HEMBB1000005	F-HEMBB1000005	0. 35
50	HEMB81000054	F-HEMBB1000054	0. 18
	HEMBB1000055	F-HEMBB1000055	0.87
	HEMBB1000144	F-HEMBB1000144	0. 47
	HEMBB1000258 HEMBB1000318	F-HEMBB1000258	0.87
55	- · -	F-HEMBB1000318	0.34
	HEMBB1000335	F-HEMBB1000335	0. 48

	HEMBB1000336	F-HEMBB1000336	0. 35
	HEMBB1000354	F-HEMBB1000354	0. 47
	HEMBB1000374	F-HEMBB1000374	0. 69
5	HEMBB1000402	F-HEMBB1000402	0. 54
	HEMBB1000404	F-HEMBB1000404	0. 47
	HEMBB1000480	F-HEMBB1000480	0. 35
	HEMBB1000493	F-HEMBB1000493	0. 33
10	HEMBB1000554	F-HEMBB1000554	0.46
	HEMBB1000573	F-HEMBB1000573	0. 34
	HEMBB1000649	F-HEMBB1000649	0. 41
	HEMBB1000652	F-HEMBB1000652	0. 46
15	HEMBB1000709	F-HEMBB1000709	0. 56
,,	HEMBB1000749	F-HEMBB1000749	0.36
	HEMBB1000790	F-HEMBB1000790,	
	HEMBB1000827	F-HEMBB1000827	0.31
	HEMBB1000831	F-HEMBB1000831	0. 42
20	HEMBB1000893	F-HEMBB1000893	0. 36
	HEMBB1001004	F-HEMBB1001004	0. 43
	HEMBB1001008	F-HEMBB1001008	0. 44
	HEMBB1001047	F-HEMBB1001047	0. 47
25	HEMBB1001119	F-HEMB81001119	0. 25
	HEMBB1001142	F-HEMBB1001142	0. 15
	HEMBB1001315	F-HEMBB1001315	0. 72
	HEMBB1001317	F-HEMBB1001317	0. 49
30	HEMBB1001326	F-HEMBB1001326	0. 44
	HEMBB1001337	F-HEMBB1001337	0.18
	HEMBB1001367	F-HEMBB1001367	0. 35
	HEMBB1001424	F-HEMBB1001424	0. 34
35	HEMBB1001436	F-HEMBB1001436	0.34
	HEMBB1001458	F-HEMBB1001458	0. 47
	HEMBB1001535	F-HEMBB1001535	0.34
	HEMBB1001536	F-HEMBB1001536	0. 22
40	HEMBB1001565	F-HEMBB1001565	0. 35
	HEMBB1001747	F-HEMBB1001747	0.40
	HEMBB1001749	F-HEMBB1001749	0. 39
	HEMBB1001797	F-HEMBB1001797	0. 79
45	HEMBB1001836	F-HEMBB1001836	0. 52
	HEMBB1001863	F-HEMBB1001863	0. 38
	HEMBB1001868	F-HEMBB1001868	0. 27
	HEMBB1001875	F-HEMBB1001875	0. 46
50	HEMBB1001911	F-HEMBB1001911	0. 87
	HEMBB1001922	F-HEMBB1001922	0. 38
	HEMBB1001925	F-HEMBB1001925	0. 33
	HEMBB1001944	F-HEMBB1001944	0. 35
55	HEMBB1001983	F-HEMBB1001983	0. 32
55	HEMBB1001996	F-HEMBB1001996	0. 39

	HEMBB1001997	F-HEMBB1001997	0. 36
	HEMBB1002045	F-HEMBB1002045	0. 20
	HEMBB1002092	F-HEMBB1002092	0. 31
5	HEMBB1002247	F-HEMBB1002247	0. 53
	HEMBB1002266	F-HEMBB1002266	0. 34
	HEMBB1002387	F-HEMBB1002387	0. 32
	HEMBB1002425	F-HEMBB1002425	0. 31
10	HEMBB1002458	F-HEMBB1002458	0. 44
	HEMBB1002522	F-HEMBB1002522	0. 56
	HEMBB1002534	F-HEMBB1002534	0. 43
	HEMBB1002579	F-HEMBB1002579	0. 21
15	HEMBB1002582	F-HEMBB1002582	0, 48
	HEMBB1002596	F-HEMBB1002596	0. 42
	HEMBB1002617	F-HEMBB1002617,	0. 47
	HEMBB1002702	F-HEMBB1002702	0. 31
20	MAMMA1000043	F-MAMMA1000043	0. 45
20	MAMMA1000092	F-MAMMA1000092	0.39
	MAMMA1000129	F-MAMMA1000129	0. 47
	MAMMA1000175	F-MAMMA1000175	0.18
or.	MAMMA1000198	F-MAMMA1000198	0.46
25	MAMMA1000221	F-MAMMA1000221	0.43
	MAMMA1000307	F-MAMMA1000307	0. 35
	MAMMA1000331	F-MAMMA1000331	0.41
	MAMMA1000356	F-MAMMA1000356	0.41
30	MAMMA1000360	F-MAMMA1000360	0. 33
	MAMMA1000402	F-MAMMA1000402	0. 35
	MAMMA1000414	F-MAMMA1000414	0. 32
	MAMMA1000444	F-MAMMA1000444	0. 13
35	MAMMA1000500	F-MAMMA1000500	0. 36
	MAMMA1000522	F-MAMMA1000522	0. 46
	MAMMA1000576	F-MAMMA1000576	0.58
	MAMMA1000594	F-MAMMA1000594	0. 69
40	MAMMA1000597	F-MAMMA1000597	0. 35
	MAMMA1000720	F-MAMMA1000720	0. 43
	MAMMA1000761	, and diding (1.000 / C)	0. 13
	MAMMA1000775	F-MAMMA1000775	0. 35
45	MAMMA1000778	F-MAMMA1000778	0. 54
	MAMMA1000798	F-MAMMA1000798	0.40
	MAMMA1000862	F-MAMMA1000862	0. 87
	MAMMA1000876	F-MAMMA1000876	0. 47
50	MAMMA1000883	F-MAMMA1000883	0.41
	MAMMA1000931	F-MAMMA1000931	0. 45
	MAMMA1000940	F-MAMMA1000940 F-MAMMA1000941	0. 35 0. 35
	MAMMA1000941 MAMMA1000943	F-MAMMA1000941	
55		• •====================================	0. 23
	MAMMA1000975	F-MAMMA1000975	0. 32

	MAMMA1001038	F-MAMMA1001038	0. 41
·	MAMMA1001162	F-MAMMA1001162	0. 18
	MAMMA1001186	F-MAMMA1001186	0. 49
5	MAMMA1001220	F-MAMMA1001220	0. 43
	MAMMA1001256	F-MAMMA1001256	0. 35
	MAMMA1001274	F-MAMMA1001274	0. 57
	MAMMA1001341	F-MAMMA1001341	0. 45
10	MAMMA1001397	F-MAMMA1001397	0.64
	MAMMA1001420	F-MAMMA1001420	0. 41
	MAMMA1001547	F-MAMMA1001547	0.39
	MAMMA1001590	F-MAMMA1001590	0.46
15	MAMMA1001670	F-MAMMA1001670	0.36
	MAMMA1001679	F-MAMMA1001679	0.46
	MAMMA1001711	F-MAMMA1001711,	0.68
	MAMMA1001745	F-MAMMA1001745	0.53
20	MAMMA1001760	F-MAMMA1001760	0.58
	MAMMA1001769	F-MAMMA1001769	0. 33
	MAMMA1001815		0. 43
	MAMMA1001820		0.19
25	MAMMA1001907	F-MAMMA1001907	0. 38
	MAMMA1002056	F-MAMMA1002056	0. 35
	MAMMA1002078	F-MAMMA1002078	0. 34
	MAMMA1002093	F-MAMMA1002093	0. 55
30	MAMMA1002125	F-MAMMA1002125	0. 39
	MAMMA1002132	F-MAMMA1002132	0.57
	MAMMA1002145	F-MAMMA1002145 F-MAMMA1002250	0.78
	MAMMA1002250 MAMMA1002311	F-MAMMA1002250	0. 72
35	MAMMA1002311	F-MAMMA1002311	0. 58 0. 19
	MAMMA1002300	F-MAMMA1002411	0. 19
	MAMMA1002411	F-MAMMA1002498	0. 37
	MAMMA1002571	F-MAMMA1002571	0. 32
40	MAMMA1002701	F-MAMMA1002701	0. 55
	MAMMA1002727	F-MAMMA1002727	0. 34
	MAMMA1002728	F-MAMMA1002728	0. 55
	MAMMA1002746		0. 45
45	MAMMA1002764	F-MAMMA1002764	0. 44
70	MAMMA1002765	F-MAMMA1002765	0. 40
	MAMMA1002820	F-MAMMA1002820	0. 32
	MAMMA1002830	F-MAMMA1002830	0. 36
50	MAMMA1002909	F-MAMMA1002909	0. 64
50	MAMMA1002941	F-MAMMA1002941	0. 42
	MAMMA1002972	F-MAMMA1002972	0. 08
	MAMMA1002973	F-MAMMA1002973	0. 35
	MAMMA1003004	F-MAMMA1003004	0. 42
55	MAMMA1003007	F-MAMMA1003007	0. 41

	MAMMA1003039	F-MAMMA1003039	0.46
	MAMMA1003089	F-MAMMA1003089	0. 58
	NT2RM4000086	F-NT2RM4000086	0.36
5	NT2RM4000265	F-NT2RM4000265_	0, 39
	NT2RM4000366	F-NT2RM4000366	0. 21
	NT2RM4000414	F-NT2RM4000414	0.35
	NT2RM4000425	F-NT2RM4000425	0. 22
10	NT2RM4000779	F-NT2RM4000779	0.43
	NT2RM4000855	F-NT2RM4000855	0.52
	NT2RM4001160	F-NT2RM4001160	0.36
	NT2RM4001313	F-NT2RM4001313	0. 38
15	NT2RM4001437	F-NT2RM4001437	0. 53
	NT2RM4001754	F-NT2RM4001754	0.35
	NT2RM4001856	F-NT2RM4001856,	0. 27
	NT2RM4001953	F-NT2RM4001953	0.59
20	NT2RM4001984	F-NT2RM4001984	0.35
	NT2RM4002390	F-NT2RM4002390	0.24
	NT2RP2000077	F-NT2RP2000077	0.35
	NT2RP2000108	F-NT2RP2000108	0.08
25	NT2RP2000183	F-NT2RP2000183	0. 43
	NT2RP2000257	F-NT2RP2000257	0.07
	NT2RP2000289	F-NT2RP2000289	0. 35
	NT2RP2000420	F-NT2RP2000420	0. 48
30	NT2RP2000678	F-NT2RP2000678	0. 55
	NT2RP2000715	F-NT2RP2000715	0. 37
	NT2RP2000842	F-NT2RP2000842	0. 40
	NT2RP2000970	F-NT2RP2000970	0. 46
35	NT2RP2001149	F-NT2RP2001149	0. 31
39	NT2RP2001226	F-NT2RP2001226	0. 44
	NT2RP2001295 NT2RP2001347	F-NT2RP2001295	0.40
	NT2RP2001347	F-NT2RP2001347	0. 41
	NT2RP2001506	F-NT2RP2001467 F-NT2RP2001506	0. 44
40	NT2RP2001569	F-NT2RP2001569	0. 17
	NT2RP2001663	F-NT2RP2001663	0. 31
	NT2RP2001936	F-NT2RP2001936	0. 37
	NT2RP2001936	F-NT2RP2001936	0. 33
45	NT2RP2002047	F-NT2RP2002047	0. 31
	NT2RP2002066	F-NT2RP2002047	0. 15 0. 11
	NT2RP2002172	F-NT2RP2002172	0. 11
	NT2RP2002172	F-NT2RP2002172	0. 54
50	NT2RP2002316	F-NT2RP2002316	0. 39
	NT2RP2002475	F-NT2RP2002475	0. 09
	NT2RP2002546	F-NT2RP2002546	0. 09
	NT2RP2002591	F-NT2RP2002591	0. 38
55	NT2RP2002643	F-NT2RP2002643	0. 55
	2002040	1 11414 2002043	J. JU

	NT2RP2002741	F-NT2RP2002741	0. 73
	NT2RP2002750	F-NT2RP2002750	0. 48
_	NT2RP2002778	F-NT2RP2002778	0.41
5	NT2RP2002857	F-NT2RP2002857	0.34
	NT2RP2003000	F-NT2RP2003000	0.56
	NT2RP2003073	F-NT2RP2003073	0.36
	NT2RP2003237	F-NT2RP2003237	0.69
10	NT2RP2003394	F-NT2RP2003394	0.34
	NT2RP2003517	F-NT2RP2003517	0.44
	NT2RP2003668	F-NT2RP2003668	0.46
	NT2RP2003988	F-NT2RP2003988	0.55
15	NT2RP2004232	F-NT2RP2004232	0.34
	NT2RP2004400	F-NT2RP2004400	0.18
	NT2RP2004512	F-NT2RP2004512,	0.11
	NT2RP2004523	F-NT2RP2004523	0.72
20	NT2RP2004587	F-NT2RP2004587	0.09
	NT2RP2004736	F-NT2RP2004736	0.70
	NT2RP2004767	F-NT2RP2004767	0.54
	NT2RP2004775	F-NT2RP2004775	0.53
25	NT2RP2004961	F-NT2RP2004961	0.49
20	NT2RP2004962	F-NT2RP2004962	0.87
	NT2RP2004982	F-NT2RP2004982	0.32
	NT2RP2005289	F-NT2RP2005289	0.24
	NT2RP2005407	F-NT2RP2005407	0.33
30	NT2RP2005531	F-NT2RP2005531	0. 20
	NT2RP2005694	F-NT2RP2005694	0. 21
	NT2RP2005726	F-NT2RP2005726	0.42
	NT2RP2005942	F-NT2RP2005942	0.10
35	NT2RP2006258	F-NT2RP2006258	0.37
	NT2RP2006261	F-NT2RP2006261	0.42
	NT2RP2006454	F-NT2RP2006454	0.46
	NT2RP2006554	F-NT2RP2006554	0.14
40	NT2RP3000055	F-NT2RP3000055	0.31
	NT2RP3000233	F-NT2RP3000233	0.54
	NT2RP3000341	F-NT2RP3000341	0.43
	NT2RP3000418		0.53
45	NT2RP3000451	F-NT2RP3000451	0. 46
	NT2RP3000561	F-NT2RP3000561	0. 47
	NT2RP3000582	F-NT2RP3000582	0. 44
	NT2RP3001007	F-NT2RP3001007	0. 26
50	NT2RP3001281	F-NT2RP3001281	0. 36
	NT2RP3001318	F-NT2RP3001318	0. 21
	NT2RP3001339	F-NT2RP3001339	0. 58
	NT2RP3001340	F-NT2RP3001340	0. 41
55	NT2RP3001383	F-NT2RP3001383	0. 36
55	NT2RP3001432	F-NT2RP3001432	0.34

	NT2RP3001580	F-NT2RP3001580	0. 45
	NT2RP3001589	F-NT2RP3001589	0. 48
	NT2RP3001898	F-NT2RP3001898	0.19
5	NT2RP3002004	F-NT2RP3002004	0.36
	NT2RP3002173	F-NT2RP3002173	0.39
	NT2RP3003133	F-NT2RP3003133	0. 35
	NT2RP3003264	F-NT2RP3003264	0.13
10	NT2RP3003346	F-NT2RP3003346	0. 72
	NT2RP3003403	F-NT2RP3003403	0. 45
	NT2RP3003433	F-NT2RP3003433	0. 24
	NT2RP3003576	F-NT2RP3003576	0.35
15	NT2RP3003625	F-NT2RP3003625	0. 43
	NT2RP3003665	F-NT2RP3003665	0.32
	NT2RP3003800	F-NT2RP3003800,	0.34
	NT2RP3003828	F-NT2RP3003828	0. 52
20	NT2RP3003842	F-NT2RP3003842	0. 23
20	NT2RP3004070	F-NT2RP3004070	0. 32
	NT2RP3004470	F-NT2RP3004470	0.46
	NT2RP4000023	F-NT2RP4000023	0. 33
25	NT2RP4000035	F-NT2RP4000035	0. 56
25	NT2RP4000102	F-NT2RP4000102	0. 38
	NT2RP4000167	F-NT2RP4000167	0. 42
	NT2RP4000214	F-NT2RP4000214	0. 33
20	NT2RP4000218	F-NT2RP4000218	0. 37
30	NT2RP4000424	F-NT2RP4000424	0. 48
	NT2RP4000915	F-NT2RP4000915	0. 37
	NT2RP4002075	F-NT2RP4002075	0. 43
0.5	0VARC1000017	F-0VARC1000017	0. 14
35	OVARC1000068 OVARC1000085	F-0VARC1000068 F-0VARC1000085	0. 17
	0VARC1000085	F-0VARC1000085	0. 46 0. 58
	0VARC1000092	F-0VARC1000092	0. 58
	OVARC1000143	F-0VARC1000143	0. 40
40	OVARC1000414	F-0VARC1000414	0. 15
	OVARC1000496	F-0VARC1000496	0. 36
	0VARC1000526	F-0VARC1000526	0. 33
	0VARC1000948	F-0VARC1000948	0. 71
45	0VARC1001011	F-0VARC1001011	0. 42
	OVARC1001240	F-0VARC1001240	0. 14
	0VARC1001600	F-0VARC1001600	0. 37
	0VARC1001805	F-0VARC1001805	0. 51
50	0VARC1001813	F-0VARC1001813	0. 58
	OVARC1001846	F-0VARC1001846	0. 59
	PLACE1000540	F-PLACE1000540	0. 35
	PLACE1000599	F-PLACE1000599	0. 37
55	PLACE1001088	F-PLACE1001088	0.36

	PLACE1001323	F-PLACE1001323	0. 21
	PLACE1001377	F-PLACE1001377	0. 41
	PLACE1001440	F-PLACE1001440	0.45
5	PLACE1001517	F-PLACE1001517	
	PLACE1001672	F-PLACE1001672	0.35
	PLACE1001705	F-PLACE1001705	0. 33
	PLACE1001756	F-PLACE1001756	0.67
10	PLACE1001750	F-PLACE1002157	0.68
	PLACE1002157	F-PLACE1002137	0.55
	PLACE1002203	F-PLACE1002227	0. 33
	PLACE1002259	F-PLACE1002259	0. 14
	PLACE1002239	F-PLACE1002239	0.47
15	PLACE1002319 PLACE1002399		
		F-PLACE1002399	0.31
	PLACE1002477	F-PLACE1002477,	
	PLACE1002500	F-PLACE1002500	0. 26
20	PLACE1002583	F-PLACE1002583	0.49
	PLACE1002604	F-PLACE1002604	0. 25
	PLACE1002772	F-PLACE1002772	0.13
	PLACE1002853		0. 46
25	PLACE1002968	F-PLACE1002968	0. 33
	PLACE1003238	F-PLACE1003238	0. 46
	PLACE1003420	F-PLACE1003420	0. 42
	PLACE1003478	F-PLACE1003478	0. 20
•	PLACE1003566	F-PLACE1003566	0. 35
30	PLACE1003593	F-PLACE1003593	0. 42
	PLACE1003618	F-PLACE1003618	0.36
	PLACE1004274	F-PLACE1004274	0.41
	PLACE1004681	F-PLACE1004681	0. 25
35	PLACE1004716	F-PLACE1004716	0. 49
	PLACE1004773	F-PLACE1004773	0. 31
	PLACE1004815	F-PLACE1004815	0. 72
	PLACE1004836	F-PLACE1004836	0. 39
40	PLACE1004913	F-PLACE1004913	0. 44
	PLACE1004979	F-PLACE1004979	0.44
	PLACE1005052	F-PLACE1005052	0. 63
	PLACE1005086	F-PLACE1005086	0. 57
45	PLACE1005108	F-PLACE1005108	0. 17
	PLACE1005128	F-PLACE1005128	0. 40
	PLACE1005176	F-PLACE1005176	0. 48
	PLACE1005467	F-PLACE1005467	0. 37
	PLACE1005639	F-PLACE1005639	0. 33
50	PLACE1005850	F-PLACE1005850	0. 54
	PLACE1005932	F-PLACE1005932	0. 20
	PLACE1006003	F-PLACE1006003	0.71
	PLACE1006017	F-PLACE1006017	0. 42
55	PLACE1006288	F-PLACE1006288	0. 34

PLACE1006368 F-PLACE1006368 0. 10		PLACE1006318	F-PLACE1006318	0. 22
5 PLACE1006506 P-PLACE1006506 P-PLACE1006629 0. 28 PLACE1006629 P-PLACE1006629 0. 44 PLACE1006795 P-PLACE1006904 0. 20 0. 30 PLACE1006904 P-PLACE1006904 0. 20 0. 30 PLACE1006904 P-PLACE1007478 0. 35 PLACE1007478 P-PLACE1007478 0. 70 0. 20 0. 20 PLACE1007478 10 PLACE1007557 P-PLACE1007478 P-PLACE1007478 P-PLACE1007473 P-PLACE1007473 P-PLACE1007477 P-PLACE1007877 P-PLACE1007877 P-PLACE1007877 P-PLACE1007877 P-PLACE1007877 P-PLACE1007877 P-PLACE1008181 P-PLACE1008181 P-PLACE1008181 P-PLACE1008330 PLACE1008330 PLACE1008330 P-PLACE1008330 P-PLACE1008330 P-PLACE1008330 P-PLACE1008330 P-PLACE1008330 P-PLACE1008330 P-PLACE1008331 PLACE1008631 P-PLACE1008631 P-PLACE1008631 P-PLACE1008631 P-PLACE1008631 P-PLACE1008631 P-PLACE1008631 P-PLACE1009048 P-PLACE1009048 P-PLACE1009048 P-PLACE1009048 P-PLACE1009049 PLACE1009048 P-PLACE1009049 PLACE1009049 P-PLACE1009049 P-PLACE1010562 P-PLACE10110802 P-PLACE1011185 P-PLACE1011		PLACE1006368	F-PLACE1006368	0.10
5 PLACE1006506 P-PLACE1006506 PLACE1006629 0. 28 PLACE1006629 PLACE1006629 0. 44 PLACE1006795 P-PLACE1006904 0. 20 0. 44 PLACE1006904 P-PLACE1006904 0. 20 0		PLACE1006371	F-PLACE1006371	
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55 PLACE2000061 F-PLACE2000061 0.34				
	55	PLAUE2000061	F-PLACE2000061	0. 34

	PLACE2000187	F-PLACE2000187	0.40
	PLACE2000216	F-PLACE2000187	0. 40
	PLACE2000335	F-PLACE2000216	0. 32
5	PLACE2000347		0. 78
•	PLACE2000347	F-PLACE2000347	0.36
		F-PLACE2000366	0. 33
	PLACE2000394	F-PLACE2000394	0. 54
	PLACE2000398	F-PLACE2000398	0. 44
10	PLACE2000425	F-PLACE2000425	0. 47
	PLACE2000450	F-PLACE2000450	0. 53
	PLACE2000477	F-PLACE2000477	0. 36
	PLACE3000119	F-PLACE3000119	0. 40
15	PLACE3000207	F-PLACE3000207	0. 32
	PLACE3000230	F-PLACE3000230	0. 51
	PLACE3000271	F-PLACE3000271,	
	PLACE3000373	F-PLACE3000373	0. 37
20	PLACE3000399	F-PLACE3000399	0. 42
	PLACE3000401	F-PLACE3000401	0. 37
	PLACE3000406	F-PLACE3000406	0. 46
	PLACE4000247	F-PLACE4000247	0. 64
25	PLACE4000320	F-PLACE4000320	0. 36
20	PLACE4000344	F-PLACE4000344	0. 31
	PLACE4000367	F-PLACE4000367	0. 32
	PLACE4000401	F-PLACE4000401	0. 54
	PLACE4000548	F-PLACE4000548	0. 26
30	THYR01000111	F-THYR01000111	0. 53
	THYR01000129	F-THYR01000129	0. 45
	THYR01000187	F-THYR01000187	0. 46
	THYR01000279	F-THYR01000279	0.09
35	THYR01000484	F-THYR01000484	0. 36
	THYR01000596	F-THYR01000596	0. 47
	THYR01000625	F-THYR01000625	0.59
	THYR01000793	F-THYR01000793	0. 17
40	THYR01000815	F-THYR01000815	0. 58
	THYR01000865	F-THYR01000865	0. 55
	THYR01001003	F-THYR01001003	0. 34
	THYR01001031	F-THYR01001031	0. 43
45	THYR01001133	F-THYR01001133	0. 52
	THYR01001321	F-THYR01001321	0. 37
	THYR01001401	F-THYR01001401	0. 36
	THYR01001426	F-THYR01001426	0. 32
		F-THYR01001434	0. 39
50	THYR01001559	F-THYR01001559	0. 45
	THYR01001570	F-THYR01001570	0. 35
	THYR01001706	F-THYR01001706	0. 45
	THYR01001746	F-THYR01001746	0. 43
55	THYR01001772	F-THYR01001772	0. 37

THYR01001907	F-THYR01001907	0. 33
Y79AA1000033	F-Y79AA1000033	0. 53
Y79AA1000346	F-Y79AA1000346	0. 53
Y79AA1000410	F-Y79AA1000410	0.13
Y79AA1000805	F-Y79AA1000805	0.32
Y79AA1001692	F-Y79AA1001692	0. 45
Y79AA1002103	F-Y79AA1002103	0. 15
Y79AA1002220	F-Y79AA1002220	0.49

15 EXAMPLE 13

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Full-length sequence analysis and homology search

[0192] Full-length sequence was determined for each selected cDNA clones. The nucleotide sequence determination was performed mainly by the dye-terminator method using custom synthesized DNA primers according to the primer walking procedure (custom synthesized DNA primers were used for sequencing; sequencing reaction was performed with DNA sequencing reagent supplied by PE Biosystems according to the supplier's manual; and the samples were analyzed in an automatic sequencer made by the same supplier). Sequence determination of some clones was carried out in the same manner but using a Licor DNA sequencer. Overlapping partial nucleotide sequences, which were obtained by the above-described method, were assembled together to determine a full-length nucleotide sequence. Amino acid sequences were then deduced from the determined full-length nucleotide sequences. However, amino acid sequence is not shown for a clone of which coding region was hard to be deduced or of which amino acid sequence has less than 100 amino acid residues. SEQ ID NOs corresponding to the respective clones are indicated in Table 350 and Table 351.

[0193] GenBank, Swiss-Prot and UniGene were searched for the determined nucleotide sequences by BLAST analysis. Matching data of cDNA clone which exhibits higher homology and of which functions are easily predicted based on the nucleotide sequences and the deduced amino acid sequences are selected from the BLAST analysis matching data with P value of 10⁻⁴ or less. The matching data selected are listed herein. The results of homology search 6, 12, 13, and 14 are indicated in the last part of this specification. However, there are some clones that did not match the criteria for judgment and such matching data of BLAST analysis are not shown herein.

EXAMPLE 14

Novel full-length cDNA clone obtained from a cDNA library prepared by oligo-capping method

[0194] A cDNA clone, NT2RP4002298, was obtained from a cDNA library, NT2RP4 (see Example 1), prepared by oligo-capping method. Analysis of the entire nucleotide sequence of the clone has revealed that the clone encodes a novel protein consisting of 775 amino acids. The ATGpr1 score at the initiation codon of the amino acid sequence is 0.16, and therefore, the fullness ratio is low. However, the sequence can be full-length.

[0195] The full-length nucleotide sequence of NT2RP4002298 is shown in SEQ ID NO: 12370, and the deduced amino acid sequence encoded by the clone NT2RP4002298 is shown in SEQ ID NO: 12371.

EXAMPLE 15

50 Gene expression analysis with hybridization using high density DNA filter

[0196] Nylon membrane for DNA spotting was prepared according to the following procedure. E. coli was cultured in each well of a 96-well plate (in a LB medium at 37°C for 16 hours). A small aliquot of each culture was suspended in 10 µl of sterile water in a well of a 96-well plate. The plate was heated at 100°C for 10 minutes. Then the boiled samples were analyzed by PCR reaction. PCR was performed in a 20 µl solution by using TaKaRa PCR Amplification Kit (Takara) according to the supplier's protocol. Primers used for the amplification of an insert cDNA in a plasmid were a pair of sequencing primers, ME761FW (5' tacggaagtgttacttctgc 3' / SEQ ID NO: 13290) and ME1250RV (5' tgtgggaggttttttctcta 3' / SEQ ID NO: 13291), or a pair of primers, M13M4 (5' gttttcccagtcacgac 3' / SEQ ID NO: 13292)

and M13RV (5' caggaaacagctatgac 3' / SEQ ID NO: 13293). PCR reaction was performed in a thermal cycler, GeneAmp System 9600 (PE Biosystems). The cycling profile consisted of pre-heat at 95°C for 5 mimutes; 10 cycles of denaturation at 95°C for 10 seconds, and annealing/extension at 68°C for 1 minute; 20 cycles of denaturation at 98°C for 20 seconds and annealing/extension at 60°C for 3 minutes; and final extension at 72°C for 10 minutes. After the PCR reaction, the 20 µl reaction solution was loaded onto a 1 % agarose gel and fractionated by electrophoresis. DNA on the gel was stained with ethicium bromide to confirm the amplification of cDNA. When cDNAs were barely amplified by PCR, plasmids containing the corresponding insert cDNAs were prepared by the alkali-extraction method (J. Sambrook, E. F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

[0197] Preparation of DNA array was carried out by the following procedure. An Aliquot of a DNA solution was added in each well of a 384-well plate. DNA was spotted onto a nylon membrane (Boehringer) by using a 384-pin tool of Biomek 2000 Laboratory Automation Sysytem (Beckman-Coulter). Specifically, the 384-well plate containing the DNA was placed under the 384-pin tool. The independent 384 needles were simultaneously dipped into the DNA solution for DNA deposition. The needles were gently pressed onto a nylon membrane and the DNA deposited at the tips of needles was spotted onto the membrane. Denaturation of the spotted DNA and immobilization of the DNA on the nylon membrane were carried out according to usual manners (J. Sambrook, E.F., Fritsh, & T. Maniatis, "Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989).

[0198] Hybridization probe used was radioisotope-labeled 1st strand cDNA. The 1st strand cDNA synthesis was performed by using Thermoscript(TM) RT-PCR System (GIBCO). Specifically, the 1st strand cDNA was synthesized by using 1.5 μ g mRNAs from various human tissues (Clontech), 1 μ l aliquots of 50 μ M Oligo(dT)20 and 50 μ Ci [α ³³P] dATP according to an attached protocol. Probe purification was carried out by using ProbeQuant(TM) G-50 micro column (Amersham-Pharmacia Biotech) according to an attached protocol. In the next step, 2 units of E. coli RNasH were added to the reaction mixture. The mixture was incubated at room temperature for 10 minutes and then 100 μ g of human COT-1 DNA (GIBCO) was added thereto. The mixture was incubated at 97°C for 10 minutes and then was allowed to stand on ice to give hybridization probe.

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[0199] Hybridization of the radioisotope-labeled probe to the DNA array was performed in a usual manner (J. Sambrook, E.F., Fritsh, & T. Maniatis, Molecular Cloning, A laboratory manual/ 2nd edition, Cold Spring Harbor Laboratory Press, 1989). The membrane was washed as follows: the nylon membrane was incubated in Washing solution 1 (2 x SSC, 1% SDS) at room temperature (about 26°C) for 20 minutes and this washing was repeated 3 times; then the membrane was washed 3 times by incubating it in Washing solution 2 (0.1 × SSC, 1% SDS) at 65°C for 20 minutes. Autoradiography was performed by using an image plate for BAS2000 (Fuji Photo Film Co., Ltd.). Specifically, the nylon membrane with probe hybridized thereon was wrapped with a piece of Saran Wrap and brought into tight contact with the image plate on the light-sensitive surface. The membrane with the image plate was placed in an imaging cassette for radioisotope and allowed to stand in dark place for 4 hours. The radioactivity recorded on the image plate was analyzed by using BAS2000 (Fuji Photo Film Co., Ltd.). The activity was subjected to electronic conversion and recorded as an image file of autoradiogram. The signal intensity of each DNA spot was analyzed by using Visage High Density Grid Analysis Systems (Genomic Solutions Inc.). The signal intensity was converted into numerical data. The data were taken by duplicated measurements. The reproducibility was assessed by comparing the signal intensities of the corresponding spots on the duplicated DNA filters that were hybridized to a single DNA probe (Figure 2). The ratio between the corresponding spots falls within a range of 2 or less in 95% of entire spots and the correlation coefficient is r=0.97. Thus the reproducibility is assumed to be satisfactory.

[0200] The detection sensitivity in gene expression analysis was estimated by examining increases in the signal intensity of probe concentration-dependent spot in hybridization using a probe complementary to the DNA spotted on the nylon membrane. DNA used was PLACE1008092 (same as DNA deposited in GenBank under an Accession No. AF107253). The DNA array with DNA of PLACE1008092 was prepared according to the above-mentioned method. The probe used was prepared as follows: mRNA was synthesized in vitro from the clone, PLACE1008092. By using this mRNA as a template, radioisotope-labeled 1st strand cDNA was synthesized in the same manner as described above, and the cDNA was used as the probe. The cDNA PLACE1008092 was inserted into pBluescript SK(-), of which T7 promoter was ligated to the 5' end of the cDNA, to give a plasmid for in vitro synthesis of the mRNA from PLACE1008092. Specifically, the PLACE1008092 insert was cut out from pME18SFL3 carrying the cDNA at a DraIII site thereof by Xhol digestion. The resulting PLACE1008092 fragment was ligated to Xhol-predigested pBluescript SK $\,$ (-) by using DNA ligation kit ver.2 (Takara). The in-vitro mRNA synthesis from PLACE1008092 inserted in pBluescript SK(-) was carried out by using Ampliscribe(TM) T7 high yield transcription kit (Epicentre technologies). Hybridization and the analysis of signal intensity of each DNA spot were conducted by the same methods as described above. When the probe concentration is $1 \times 10^7 \, \mu \text{g/ml}$ or less, there was no increase of signal intensity proportional to the probe concentration. Therefore it was assumed to be difficult to compare the signals with one another in the concentration range. Thus the spots with the intensity of 40 or less were indiscriminately taken as low-level signals (Figure 3). Within a concentration of the probe ranging from $1 \times 10^7 \,\mu\text{g/ml}$ to $0.1 \,\mu\text{g/ml}$, the signal was found to increase in a probe

concentration-dependent manner. The detection limit was then assumed to be 1:100,000 in a ratio of mRNA expression level in a sample.

[0201] Tables 12-168 (also containing clones with no description in Examples) show the expression of each cDNA in human normal tissues (heart, lung, pituitary gland, thymus, brain, kidney, liver and spleen). The expression levels are indicated by numerical values of 0-10,000. Genes that were expressed in at least a single tissue are indicated below by the corresponding clone names:

	HEMBA1000012,	HEMBA1000020,	HEMBA1000030,	HEMBA1000042,	HEMBA1000046,
10	HEMBA1000076,	HEMBA1000111,	HEMBA1000129,	HEMBA1000150,	HEMBA1000156,
	HEMBA1000158,	HEMBA1000168,	HEMBA1000185,	HEMBA1000216,	HEMBA1000227,
	HEMBA1000231,	HEMBA1000243,	HEMBA1000244,	HEMBA1000280,	HEMBA1000282,
	HEMBA1000288,	HEMBA1000303,	HEMBA1000304,	HEMBA1000327,	HEMBA1000338,
15	HEMBA1000351,	HEMBA1000355,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366,
	HEMBA1000369,	HEMBA1000376,	HEMBA1000387,	HEMBA1000392,	HEMBA1000396,
	HEMBA1000411,	HEMBA1000422,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000469,	HEMBA1000488,	HEMBA1000491,	HEMBA1000501,	HEMBA1000505,
20	HEMBA1000508,	HEMBA1000519,	HEMBA1000523,	HEMBA1000534,	HEMBA1000540,
	HEMBA1000542,	HEMBA1000545,	HEMBA1000557,	HEMBA1000561,	HEMBA1000568,
	HEMBA1000569,	HEMBA1000575,	HEMBA1000588,	HEMBA1000591,	HEMBA1000604,

	-HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000657,	HEMBA1000673,
	HEMBA1000682,	HEMBA1000702,	HEMBA1000705,	HEMBA1000726,	HEMBA1000749,
_	HEMBA1000752,	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,
5	HEMBA1000843,	HEMBA1000852,	HEMBA1000870,	HEMBA1000872,	HEMBA1000876,
	HEMBA1000908,	HEMBA1000910,	HEMBA1000918,	HEMBA1000919,	HEMBA1000934,
	HEMBA1000942,	HEMBA1000960,	HEMBA1000971,	HEMBA1000972,	HEMBA1000975,
	HEMBA1000986,	HEMBA1000991,	HEMBA1001007,	HEMBA1001008,	HEMBA1001017,
10	HEMBA1001020,	HEMBA1001051, HE	MBA1001059, HEMB	A1001060, HEMBA1	001077,
	HEMBA1001080,	HEMBA1001085,	HEMBA1001088,	HEMBA1001109,	HEMBA1001121,
	HEMBA1001123,	HEMBA1001133,	HEMBA1001137,	HEMBA1001140,	HEMBA1001172,
	HEMBA1001197,	HEMBA1001208,	HEMBA1001213,	HEMBA1001226,	HEMBA1001235,
15	HEMBA1001265,	HEMBA1001281,	HEMBA1001286,	HEMBA1001289,	HEMBA1001294,
	HEMBA1001299,	HEMBA1001302,	HEMBA1001303,	HEMBA1001310,	HEMBA1001326,
	HEMBA1001330,	HEMBA1001351,,	HEMBA1001375,	HEMBA1001377,	HEMBA1001387,
	HEMBA1001388,	HEMBA1001398,	HEMBA1001405,	HEMBA1001413,	HEMBA1001415,
20	HEMBA1001432,	HEMBA1001433,	HEMBA1001435,	HEMBA1001446,	HEMBA1001450,
	HEMBA1001454,	HEMBA1001463,	HEMBA1001476,	HEMBA1001497,	HEMBA1001510-
	HEMBA1001515,	HEMBA1001517,	HEMBA1001526,	HEMBA1001533,	HEMBA1001557,
	HEMBA1001566,	HEMBA1001569,	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,
	HEMBA1001589,	HEMBA1001595,	HEMBA1001608,	HEMBA1001620,	HEMBA1001636;
25	HEMBA1001640,	HEMBA1001647,	HEMBA1001651,	HEMBA1001655,	HEMBA1001661,
	HEMBA1001678,	HEMBA1001709,	HEMBA1001712,	HEMBA1001714,	HEMBA1001718,
	HEMBA1001731,	HEMBA1001734,	HEMBA1001745,	HEMBA1001761,	HEMBA1001781,
	HEMBA1001784,	HEMBA1001791,	HEMBA1001804,	HEMBA1001808,	HEMBA1001809,
30	HEMBA1001815,	HEMBA1001822,	HEMBA1001824,	HEMBA1001844,	HEMBA1001847,
	HEMBA1001864,	HEMBA1001866,	HEMBA1001869,	HEMBA1001888,	HEMBA1001896,
	HEMBA1001910,	HEMBA1001912,	HEMBA1001913,	HEMBA1001939,	HEMBA1001940,
	HEMBA1001964,	HEMBA1001967, HE	MBA1001987, HEMB	A1001991, HEMBA1	002003,
35	HEMBA1002008,	HEMBA1002018,	HEMBA1002035,	HEMBA1002049,	HEMBA1002092,
	HEMBA1002100,	HEMBA1002102,	HEMBA1002113,	HEMBA1002119,	HEMBA1002125,
	HEMBA1002144,	HEMBA1002150,	HEMBA1002151,	HEMBA1002160,	HEMBA1002161,
	HEMBA1002162,	HEMBA1002166,	HEMBA1002177,	HEMBA1002185,	HEMBA1002189,
40	HEMBA1002191,	HEMBA1002226,	HEMBA1002229,	HEMBA1002237,	HEMBA1002241,
	HEMBA1002265,	HEMBA1002267,	HEMBA1002270,	HEMBA1002328,	HEMBA1002337,
	HEMBA1002341,	HEMBA1002363,	HEMBA1002381,	HEMBA1002389,	HEMBA1002417,
	HEMBA1002419,	HEMBA1002439,	HEMBA1002458,	HEMBA1002462,	HEMBA1002469,
45	HEMBA1002477,	HEMBA1002486,	HEMBA1002495,	HEMBA1002498,	HEMBA1002503,
	HEMBA1002508,	HEMBA1002513,	HEMBA1002515,	HEMBA1002538,	HEMBA1002542,
	HEMBA1002552,	HEMBA1002558,	HEMBA1002561,	HEMBA1002569,	HEMBA1002590,
	HEMBA1002592,	HEMBA1002609,	HEMBA1002624,	HEMBA1002629,	HEMBA1002645,
50	HEMBA1002659,	HEMBA1002661,	HEMBA1002678,	HEMBA1002679,	HEMBA1002696,
50	HEMBA1002703,	HEMBA1002712,	HEMBA1002728,	HEMBA1002730,	HEMBA1002746,
	HEMBA1002748,	HEMBA1002750,	HEMBA1002768,	HEMBA1002777,	HEMBA1002779,
	HEMBA1002780,	HEMBA1002794,	HEMBA1002810,	HEMBA1002816,	HEMBA1002818,
	HEMBA1002826,	HEMBA1002833,	HEMBA1002863,	HEMBA1002876,	HEMBA1002896,
55	HEMBA1002921,	HEMBA1002924,	HEMBA1002934,	HEMBA1002935,	HEMBA1002944,
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	HEMBA1002968,	HEMBA1002970,	HEMBA1002971,	HEMBA1002973,	HEMBA1002997,
	HEMBA1003021,	HEMBA1003033, HEM	MBA1003034, HEMBA	41003037, HEMBA1	003041,
	HEMBA1003046,	HEMBA1003067,	HEMBA1003071,	HEMBA1003078,	HEMBA1003079,
5	HEMBA1003083,	HEMBA1003086,	HEMBA1003098,	HEMBA1003129,	HEMBA1003133,
	HEMBA1003136,	HEMBA1003142,	HEMBA1003148,	HEMBA1003166,	HEMBA1003175,
	HEMBA1003179,	HEMBA1003199,	HEMBA1003202,	HEMBA1003204,	HEMBA1003212,
	HEMBA1003220,	HEMBA1003229,	HEMBA1003235,	HEMBA1003257,	HEMBA1003273,
10	HEMBA1003276,	HEMBA1003278,	HEMBA1003281,	HEMBA1003286,	HEMBA1003296,
	HEMBA1003314,	HEMBA1003322,	HEMBA1003327,	HEMBA1003328,	HEMBA1003330,
	HEMBA1003348,	HEMBA1003370,	HEMBA1003373,	HEMBA1003376,	HEMBA1003380,
	HEMBA1003384,	HEMBA1003403,	HEMBA1003408,	HEMBA1003418,	HEMBA1003447,
15	HEMBA1003461,	HEMBA1003463,	HEMBA1003480,	HEMBA1003528,	HEMBA1003531,
	HEMBA1003538,	HEMBA1003556,	HEMBA1003571,	HEMBA1003581,	HEMBA1003591,
	HEMBA1003595,	HEMBA1003597,	HEMBA1003598,	HEMBA1003615,	HEMBA1003617,
	HEMBA1003621,	HEMBA1003637,	HEMBA1003640,	HEMBA1003645,	HEMBA1003656,
20	HEMBA1003667,	HEMBA1003680,	HEMBA1003690,	HEMBA1003692,	HEMBA1003711.
20	HEMBA1003714,	HEMBA1003715,	HEMBA1003720,	HEMBA1003725,	HEMBA1003729-
	HEMBA1003733,	HEMBA1003742,	HEMBA1003758,	HEMBA1003760,	HEMBA1003773,
	HEMBA1003803,	HEMBA1003804,	HEMBA1003805,	HEMBA1003827,	HEMBA1003836,
	HEMBA1003838,	HEMBA1003864,	HEMBA1003879,	HEMBA1003880,	HEMBA1003885;
25	HEMBA1003893,	HEMBA1003902,	HEMBA1003926,	HEMBA1003937,	HEMBA1003942,
	HEMBA1003958,	HEMBA1003978, HEM	MBA1003987, HEMBA	A1003989, HEMBA1	004011,
	HEMBA1004012,	HEMBA1004024,	HEMBA1004048,	HEMBA1004049,	HEMBA1004056,
	HEMBA1004074,	HEMBA1004086,	HEMBA1004097,	HEMBA1004111,	HEMBA1004131,
30	HEMBA1004132,	HEMBA1004133,	HEMBA1004138,	HEMBA1004146,	HEMBA1004164,
	HEMBA1004200,	HEMBA1004203,	HEMBA1004225,	HEMBA1004227,	HEMBA1004238,
	HEMBA1004246,	HEMBA1004267,	HEMBA1004274,	HEMBA1004275,	HEMBA1004289,
	HEMBA1004306,	HEMBA1004312,	HEMBA1004321,	HEMBA1004323,	HEMBA1004327,
35	HEMBA1004335,	HEMBA1004341,	HEMBA1004353,	HEMBA1004354,	HEMBA1004356,
	HEMBA1004396,	HEMBA1004405,	HEMBA1004408,	HEMBA1004429,	HEMBA1004433,
	HEMBA1004460,	HEMBA1004461,	HEMBA1004479,	HEMBA1004499,	HEMBA1004506,
	HEMBA1004507,	HEMBA1004509,	HEMBA1004534,	HEMBA1004538,	HEMBA1004542,
40	HEMBA1004554,	HEMBA1004560,	HEMBA1004577,	HEMBA1004586,	HEMBA1004596,
	HEMBA1004604,	HEMBA1004610,	HEMBA1004617,	HEMBA1004629,	HEMBA1004669,
	HEMBA1004670,	HEMBA1004672,	HEMBA1004697,	HEMBA1004705,	HEMBA1004709,
	HEMBA1004711,	HEMBA1004725,	HEMBA1004736,	HEMBA1004748,	HEMBA1004751,
45	HEMBA1004752,	HEMBA1004753,	HEMBA1004756,	HEMBA1004758,	HEMBA1004763,
	HEMBA1004771,	HEMBA1004778,	HEMBA1004803,	HEMBA1004816,	HEMBA1004847,
	HEMBA1004850,	HEMBA1004863,	HEMBA1004864,	HEMBA1004865,	HEMBA1004880,
	HEMBA1004909,	HEMBA1004918,	HEMBA1004923,	HEMBA1004930,	HEMBA1004933,
50	HEMBA1004934,	HEMBA1004944,	HEMBA1004954,	HEMBA1004960,	HEMBA1004972,
	HEMBA1004980,	HEMBA1005008, HEM	ŒA1005019, HEMB∕	11005035, HEMBA1	005039,
	HEMBA1005047,	HEMBA1005050,	HEMBA1005075,	HEMBA1005079,	HEMBA1005101,
	HEMBA1005113,	HEMBA1005123,	HEMBA1005133,	HEMBA1005149,	HEMBA1005152,
	HEMBA1005201,	HEMBA1005202,	HEMBA1005206,	HEMBA1005223,	HEMBA1005241,
55	HEMBA1005244,	HEMBA1005251,	HEMBA1005252,	HEMBA1005275,	HEMBA1005296,

	HEMBA1005304,	HEMBA1005311,	HEMBA 1005315	HEMRA1005221	UEMBA 1005220
	HEMBA1005353,	HEMBA1005359,	HEMBA1005315,	HEMBA1005331,	HEMBA 1005338,
	HEMBA1005389,		HEMBA1005367,	HEMBA1005374,	HEMBA1005382,
5	HEMBA1005443,	HEMBA1005403,	HEMBA1005408,	HEMBA1005411,	HEMBA1005423,
Ü		HEMBA1005447,	HEMBA1005468,	HEMBA1005469,	HEMBA1005472,
	HEMBA1005474,	HEMBA1005475,	HEMBA1005500,	HEMBA1005511,	HEMBA1005513,
	HEMBA1005518,	HEMBA1005520,	HEMBA1005526,	HEMBA1005530,	HEMBA1005548,
	HEMBA1005552,	HEMBA1005558,	HEMBA1005568,	HEMBA1005570,	HEMBA1005576,
10	HEMBA1005577,	HEMBA1005581,	HEMBA1005583,	HEMBA1005588,	HEMBA1005593,
	HEMBA1005595,	HEMBA1005606,	HEMBA1005609,	HEMBA1005616,	HEMBA1005627,
	HEMBA1005632,	HEMBA1005634,	HEMBA1005670,	HEMBA1005679,	HEMBA1005680,
	HEMBA1005685,	HEMBA1005705,	HEMBA1005717,	HEMBA1005732,	HEMBA1005765,
15	HEMBA1005780,	HEMBA1005813,	HEMBA1005815,	HEMBA1005822,	HEMBA1005829,
	HEMBA1005834,	HEMBA1005852,	HEMBA1005853,	HEMBA1005894,	HEMBA1005911,
	HEMBA1005921,	HEMBA1005931,	HEMBA1005934,	HEMBA1005962,	HEMBA1005990,
	HEMBA1005991,	HEMBA1005999,	HEMBA1006002,	HEMBA1006005,	HEMBA1006031,
20	HEMBA1006036,	HEMBA1006042, HE	MBA1006081, HEMB	A1006090, HEMBA1	.006091,
	HEMBA1006100,	HEMBA1006108,	HEMBA1006121,	HEMBA1006124,	HEMBA1006138,
	HEMBA1006142,	HEMBA1006155,	HEMBA1006173,	HEMBA1006182,	HEMBA1006235,
	HEMBA1006248,	HEMBA1006252,	HEMBA1006253,	HEMBA1006259,	HEMBA1006278,
25	HEMBA1006283,	HEMBA1006309,	HEMBA1006310,	HEMBA1006328,	HEMBA1006344;
20	HEMBA1006349,	HEMBA1006359,	HEMBA1006364,	HEMBA1006377,	HEMBA1006380,
	HEMBA1006381,	HEMBA1006398,	HEMBA1006416,	HEMBA1006419,	HEMBA1006421,
	HEMBA1006426,	HEMBA1006438,	HEMBA1006445,	HEMBA1006461,	HEMBA1006483,
	HEMBA1006485,	HEMBA1006486,	HEMBA1006497,	HEMBA1006502,	HEMBA1006546,
30	HEMBA1006562,	HEMBA1006569,	HEMBA1006595,	HEMBA1006597,	HEMBA1006612,
	HEMBA1006617,	HEMBA1006624,	HEMBA1006631,	HEMBA1006635,	HEMBA1006639,
	HEMBA1006643,	HEMBA1006648,	HEMBA1006652,	HEMBA1006653,	HEMBA1006659,
	HEMBA1006674,	HEMBA1006676,	HEMBA1006695,	HEMBA1006696,	HEMBA1006708,
35	HEMBA1006709,	HEMBA1006717,	HEMBA1006737,	HEMBA1006744,	HEMBA1006754,
	HEMBA1006758,	HEMBA1006779,	HEMBA1006780,	HEMBA1006795,	HEMBA1006796,
	HEMBA1006807,	HEMBA1006821,	HEMBA1006824,	HEMBA1006832,	HEMBA1006849,
	HEMBA1006865,	HEMBA1006877,	HEMBA1006885,	HEMBA1006900,	HEMBA1006914,
40	HEMBA1006921,	HEMBA1006926,	HEMBA1006936,	HEMBA1006941,	HEMBA1006949,
	HEMBA1006973,	HEMBA1006993,	HEMBA1007002,	HEMBA1007045,	HEMBA1007051,
	HEMBA1007052,	HEMBA1007062,	HEMBA1007066,	HEMBA1007073,	HEMBA1007078,
	HEMBA1007080,	HEMBA1007085, HE	MBA1007087, HEMB	A1007113, HEMBA1	•
45	HEMBA1007129,	HEMBA1007147,	HEMBA1007149,	HEMBA1007151,	HEMBA1007174,
	HEMBA1007178,	HEMBA1007194,	HEMBA1007203,	HEMBA1007206,	HEMBA1007224,
	HEMBA1007243,	HEMBA1007256,	HEMBA1007267,	HEMBA1007273,	HEMBA1007279,
	HEMBA1007288,	HEMBA1007300,	HEMBA1007301,	HEMBA1007320,	HEMBA1007322,
50	HEMBA1007327,	HEMBA1007341,	HEMBA1007347,	HEMBB1000005,	HEMBB1000008,
00	HEMBB1000018,	HEMBB1000024,	HEMBB1000025,	HEMBB1000030,	HEMBB1000036,
	HEMBB1000037,	HEMBB1000039,	HEMBB1000044,	HEMBB1000050,	HEMBB1000054,
	HEMBB1000055,	HEMBB1000059,	HEMBB1000083,	HEMBB1000089,	HEMBB1000099,
	HEMBB1000103,	HEMBB1000113,	HEMBB1000119,	HEMBB1000136,	HEMBB1000141,
55	HEMBB1000144,	HEMBB1000173,	HEMBB1000198,	HEMBB1000215,	HEMBB1000217,

	HEMBB1000218,	HEMBB1000226,	HEMBB1000240,	HEMBB1000244,	HEMBB1000258,
	HEMBB1000264,	HEMBB1000266,	HEMBB1000274,	HEMBB1000307,	HEMBB1000312,
	HEMBB1000317,	HEMBB1000318,	HEMBB1000336,	HEMBB1000337,	HEMBB1000338.
5	HEMBB1000339,	HEMBB1000341,	HEMBB1000343.	HEMBB1000354,	HEMBB1000369.
	HEMBB1000374,	HEMBB1000376,	HEMBB1000391,	HEMBB1000402,	HEMBB1000420.
	HEMBB1000434,	HEMBB1000438,	HEMBB1000441,	HEMBB1000455,	HEMBB1000472,
	HEMBB1000480,	HEMBB1000490,	HEMBB1000491,	HEMBB1000493,	HEMBB1000510,
10	HEMBB1000523,	HEMBB1000530,	HEMBB1000550,	HEMBB1000554,	HEMBB1000556,
	HEMBB1000564,	HEMBB1000573,	HEMBB1000575,	HEMBB1000586,	HEMBB1000589,
	HEMBB1000591,	HEMBB1000593, HE	MBB1000598, HEME	BB1000623, HEMBB1	•
	HEMBB1000631,	HEMBB1000632,	HEMBB1000637,	HEMBB1000638,	HEMBB1000643,
15	HEMBB1000649,	HEMBB1000652,	HEMBB1000671,	HEMBB1000684,	HEMBB1000693,
	HEMBB1000705,	HEMBB1000706,	HEMBB1000709,	HEMBB1000726,	HEMBB1000738,
	HEMBB1000749,	HEMBB1000763,	HEMBB1000770,	HEMBB1000781,	HEMBB1000790,
	HEMBB1000807,	HEMBB1000810,	HEMBB1000821,	HEMBB1000826,	HEMBB1000827,
20	HEMBB1000835,	HEMBB1000840,	HEMBB1000848,	HEMBB1000870,	HEMBB1000887,
	HEMBB1000890,	HEMBB1000893,	HEMBB1000908,	HEMBB1000910,	HEMBB1000913,
	HEMBB1000915,	HEMBB1000917,	HEMBB1000927,	HEMBB1000947,	HEMBB1000959,
	HEMBB1000975,	HEMBB1000991,	HEMBB1000996,	HEMBB1001014,	HEMBB1001020,
25	HEMBB1001024,	HEMBB1001037,	HEMBB1001047,	HEMBB1001056,	HEMBB1001058;
25	HEMBB1001060,	HEMBB1001063,	HEMBB1001068,	HEMBB1001096,	HEMBB1001102,
	HEMBB1001105,	HEMBB1001112,	HEMBB1001114,	HEMBB1001119,	HEMBB1001126,
	HEMBB1001133,	HEMBB1001137,	HEMBB1001142,	HEMBB1001151,	HEMBB1001153,
	HEMBB1001169,	HEMBB1001175,	HEMBB1001177,	HEMBB1001182,	HEMBB1001208,
30	HEMBB1001209,	HEMBB1001210,	HEMBB1001218,	HEMBB1001234,	HEMBB1001249,
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40	Y79AA1001603,	Y79AA1001613,	Y79AA1001647,	Y79AA1001665,	Y79AA1001679,
40	Y79AA1001692,	Y79AA1001705,	Y79AA1001711,	Y79AA1001805,	Y79AA1001827,
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45	Y79AA1002211, Y79AA1002351,	Y79AA1002229, Y79AA1002361.	Y79AA1002234,	Y79AA1002246, Y79AA1002433,	Y79AA1002258,
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[0202] Genes that were expressed in all the tissues tested are indicated below by the corresponding clone names:

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	HEMBA1000508,	HEMBA1000519,	HEMBA1000575,	HEMBA1000655,	HEMBA1000682,
	HEMBA1000702,	HEMBA1000726,	HEMBA1000774,	HEMBA1000960,	HEMBA1000986,
5	HEMBA1001051,	HEMBA1001085,	HEMBA1001109,	HEMBA1001197,	HEMBA1001226,
	HEMBA1001286,	HEMBA1001299,	HEMBA1001377,	HEMBA1001435,	HEMBA1001476,
	HEMBA1001510,	HEMBA1001557,	HEMBA1001570,	HEMBA1001581,	HEMBA1001651,
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10	HEMBA1002102,	HEMBA1002113,	HEMBA1002150,	HEMBA1002160,	HEMBA1002166,

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	i modiova018,	PLACE1002655, PL	nceluvalia, PLAC	EIUUZOOI, PLACEI	vu3200,

	PLACE1003256,	PLACE1003361,	PLACE1003493,	·PLACE1003519,	PLACE1003528,
	PLACE1003566,	PLACE1003592,	PLACE1003870,	PLACE1003968,	PLACE1004103,
5	PLACE1004149,	PLACE1004156,	PLACE1004161,	PLACE1004242,	PLACE1004336,
· ·	PLACE1004358,	PLACE1004672,	PLACE1004736,	PLACE1004824,	PLACE1004900,
	PLACE1004979,	PLACE1005086,	PLACE1005101,	PLACE1005102,	PLACE1005128,
	PLACE1005232,	PLACE1005528,	PLACE1006002,	PLACE1006139,	PLACE1006412,
10	PLACE1006540,	PLACE1007132,	PLACE1007367,	PLACE1008405,	PLACE1008457,
	PLACE1010310,	PLACE1011056,	PLACE1011340,	PLACE1011646,	PLACE1011783,
	PLACE2000003,	PLACE2000030,	PLACE2000039,	PLACE2000047,	PLACE2000124,
	PLACE2000235,	PLACE2000305,	PLACE2000335,	PLACE2000347,	PLACE2000411.
15	PLACE2000419,	PLACE2000435,	PLACE2000450,	PLACE2000465,	PLACE3000004,
	PLACE3000009,	PLACE3000070,	PLACE3000124,	PLACE3000136,	PLACE3000145,
	PLACE3000155,	PLACE3000158,	PLACE3000207,	PLACE3000254,	PLACE3000271,
	PLACE3000304,	PLACE3000331,	PLACE3000399,	PLACE3000401,	PLACE3000455,
20	PLACE3000475,	PLACE4000009,	PLACE4000049,	PLACE4000128,	PLACE4000131,
	PLACE4000192,	PLACE4000211,	PLACE4000250,	PLACE4000323,	PLACE4000445,
	PLACE4000450,	PLACE4000465,	PLACE4000612,	THYR01000085,	THYR01000132,
	THYR01000186,	THYR01000484,	THYR01000569,	THYR01000699,	THYR01000712,
25	THYR01000815,	THYRO1001173,	THYR01001189,	THYR01001401,	THYR01001406,
	THYR01001411,	THYR01001426, THY	YR01001480, THYF	R01001487, THYR01	001537,
	THYR01001637,	THYRO1001772,	THYR01001793,	THYR01001828,	THYR01001854,
	Y79AA1000059,	Y79AA1000131,	Y79AA1000202,	Y79AA1000214,	Y79AA1000231,
30	Y79AA1000313,	Y79AA1000342,	Y79AA1000349,	Y79AA1000410,	Y79AA1000539,
	Y79AA1000560,	Y79AA1000589,	Y79AA1000833,	Y79AA1000985,	Y79AA1001077,
	Y79AA1001145,	Y79AA1001216,	Y79AA1001228,	Y79AA1001299,	Y79AA1001402,
	Y79AA1001548,	Y79AA1001603,	Y79AA1001613,	Y79AA1001805,	Y79AA1002472,
35	Y79AA1002482.				

[0203] Genes that were expressed at low levels in any of the tissues tested are indicated below by the corresponding clone names:

		HEMBA1000005,	HEMBA1000050,	HEMBA1000141,	HEMBA1000180,	HEMBA1000193,
		HEMBA1000201,	HEMBA1000213,	HEMBA1000251,	HEMBA1000264,	HEMBA1000290,
		HEMBA1000302,	HEMBA1000307,	HEMBA1000333,	HEMBA1000390,	HEMBA1000418,
	5	HEMBA1000434,	HEMBA1000442,	HEMBA1000460,	HEMBA1000464,	HEMBA1000490,
		HEMBA1000504,	HEMBA1000518,	HEMBA1000520,	HEMBA1000531,	HEMBA1000555,
		HEMBA1000563,	HEMBA1000592,	HEMBA1000594,	HEMBA1000608,	HEMBA1000637,
		HEMBA1000662,	HEMBA1000686,	HEMBA1000719,	HEMBA1000722,	HEMBA1000727,
1	0	HEMBA1000747,	HEMBA1000773,	HEMBA1000817,	HEMBA1000822,	HEMBA1000851,
		HEMBA1000867,	HEMBA1000869,	HEMBA1000943, ~	HEMBA1000946,	HEMBA1000968,
		HEMBA1000974,	HEMBA1000985,	HEMBA1001009,	HEMBA1001019,	HEMBA1001022,
		HEMBA1001024,	HEMBA1001026,	HEMBA1001043,	HEMBA1001052,	HEMBA1001071,
1	5	HEMBA1001094,	HEMBA1001099,	HEMBA1001122,	HEMBA1001174,	HEMBA1001247,
		HEMBA1001257,	HEMBA1001319,	HEMBA1001323,	HEMBA1001327,	HEMBA1001361,
		HEMBA1001383,	HEMBA1001391,	HEMBA1001407,	HEMBA1001411,	HEMBA1001442,
		HEMBA1001455,	HEMBA1001478,	HEMBA1001522,	HEMBA1001585,	HEMBA1001635,
2	0	HEMBA1001658,	HEMBA1001672,	HEMBA1001675,	HEMBA1001681,	HEMBA1001702,
		HEMBA1001711,	HEMBA1001723,	HEMBA1001744,	HEMBA1001746,	HEMBA1001800,

	HEMBA1001803,	HEMBA1001819,	HEMBA1001820,	HEMBA1001835,	HEMBA1001861.
	HEMBA1001915,	HEMBA1001918,	HEMBA1001921,	HEMBA1001942,	HEMBA1001945,
	HEMBA1001950,	HEMBA1001960, HE	MBA1001962, HEMB	A1001979, HEMBA1	•
5	HEMBA1002039,	HEMBA1002084,	HEMBA1002139,	HEMBA1002153,	HEMBA1002199,
	HEMBA1002204,	HEMBA1002212,	HEMBA1002215,	HEMBA1002253,	HEMBA1002257,
	HEMBA1002321,	HEMBA1002348,	HEMBA1002349,	HEMBA1002430,	HEMBA1002460,
	HEMBA1002475,	HEMBA1002547,	HEMBA1002555,	HEMBA1002583,	HEMBA1002621,
10	HEMBA1002628,	HEMBA1002651,	HEMBA1002666,	HEMBA1002688,	HEMBA1002716,
	HEMBA1002742,	HEMBA1002770,	HEMBA1002801,	HEMBA1002850,	HEMBA1002886,
	HEMBA1002937,	HEMBA1002939,	HEMBA1002951,	HEMBA1002954,	HEMBA1002999,
	HEMBA1003035,	HEMBA1003064,	HEMBA1003077,	HEMBA1003096,	HEMBA1003117,
15	HEMBA1003197,	HEMBA1003222,	HEMBA1003250,	HEMBA1003291,	HEMBA1003304,
	HEMBA1003309,	HEMBA1003369,	HEMBA1003395,	HEMBA1003402,	HEMBA1003417,
	HEMBA1003433,	HEMBA1003545,	HEMBA1003548,	HEMBA1003555,	HEMBA1003560,
	HEMBA1003568,	HEMBA1003569,	HEMBA1003579,	HEMBA1003622,	HEMBA1003630,
20	HEMBA1003646,	HEMBA1003662,	HEMBA1003679,	HEMBA1003684,	HEMBA1003783,
	HEMBA1003784,	HEMBA1003799,	HEMBA1003807,	HEMBA1003856,	HEMBA1003866,
	HEMBA1003908,	HEMBA1003939,	HEMBA1003950,	HEMBA1003953,	HEMBA1003959,
	HEMBA1003976,	HEMBA1003985,	HEMBA1004000,	HEMBA1004015,	HEMBA1004038,
25	HEMBA1004042,	HEMBA1004045,	HEMBA1004055,	HEMBA1004143,	HEMBA1004150,
25	HEMBA1004168,	HEMBA1004199,	HEMBA1004202,	HEMBA1004207,	HEMBA1004241,
	HEMBA1004248,	HEMBA1004264,	HEMBA1004272,	HEMBA1004276,	HEMBA1004286,
	HEMBA1004295,	HEMBA1004330, HE	MBA1004334, HEMB	A1004366, HEMBA1	004372,
	HEMBA1004389,	HEMBA1004394,	HEMBA1004482,	HEMBA1004502,	HEMBA1004573,
30	HEMBA1004631,	HEMBA1004632,	HEMBA1004637,	HEMBA1004638,	HEMBA1004666,
	HEMBA1004693,	HEMBA1004730,	HEMBA1004733,	HEMBA1004734,	HEMBA1004768,
	HEMBA1004770,	HEMBA1004776,	HEMBA1004795,	HEMBA1004806,	HEMBA1004807,
	HEMBA1004820,	HEMBA1004889,	HEMBA1004900,	HEMBA1004929,	HEMBA1004956,
35	HEMBA1004973,	HEMBA1004977,	HEMBA1004978,	HEMBA1004983,	HEMBA1004995,
	HEMBA1005009,	HEMBA1005029,	HEMBA1005062,	HEMBA1005066,	HEMBA1005083,
	HEMBA1005159,	HEMBA1005185,	HEMBA1005219,	HEMBA1005232,	HEMBA1005274,
	HEMBA1005293,	HEMBA1005314,	HEMBA1005318,	HEMBA1005372,	HEMBA1005394,
40	HEMBA1005410,	HEMBA1005426,	HEMBA1005497,	HEMBA1005506,	HEMBA1005508,
•	HEMBA1005517,	HEMBA1005528,	HEMBA1005582,	HEMBA1005621,	HEMBA1005631,
	HEMBA1005666,	HEMBA1005699,	HEMBA1005737,	HEMBA1005746,	HEMBA1005755,
	HEMBA1005884,	HEMBA1005891,	HEMBA1005909,	HEMBA1005963,	HEMBA1006035,
45	HEMBA1006067,	HEMBA1006130,	HEMBA1006158,	HEMBA1006198,	HEMBA1006268,
	HEMBA1006272,	HEMBA1006284,	HEMBA1006291,	HEMBA1006293,	HEMBA1006334,
	HEMBA1006347,	HEMBA1006424,	HEMBA1006446,	HEMBA1006467,	HEMBA1006471,
	HEMBA1006474,	HEMBA1006489,	HEMBA1006492,	HEMBA1006494,	HEMBA1006507,
50	HEMBA1006521,	HEMBA1006530,	HEMBA1006535,	HEMBA1006540,	HEMBA1006559,
	HEMBA1006566,	HEMBA1006579,	HEMBA1006583,	HEMBA1006665,	HEMBA1006682,
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	HEMBA1006996,	HEMBA1007017,	HEMBA1007018,	HEMBA1007112,	HEMBA1007251,
55	HEMBA1007281,	HEMBA1007319,	HEMBA1007342,	HEMBB1000048,	HEMBB1000175,
	HEMBB1000250,	HEMBB1000272,	HEMBB1000284,	HEMBB1000335,	HEMBB1000399,

	HEMBB1000404,	HEMBB1000449,	HEMBB1000487,	HEMBB1000518,	HEMBB1000592.
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	HEMBB1000794,	HEMBB1000822,	HEMBB1000831,	HEMBB1000852,	HEMBB1000876,
5	HEMBB1000883,	HEMBB1000888,	HEMBB1000973,	HEMBB1000981.	HEMBB1000985,
	HEMBB1001004,	HEMBB1001008.	HEMBB1001011.	HEMBB1001051,	HEMBB1001117,
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10	HEMBB1001356,	HEMBB1001364,	HEMBB1001384,	HEMBB1001387,	HEMBB1001394,
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15	HEMBB1001760,	HEMBB1001762,	HEMBB1001785,	HEMBB1001797,	HEMBB1001839.
	HEMBB1001867,	HEMBB1001868,	HEMBB1001872,	HEMBB1001874,	HEMBB1001875,
	HEMBB1001899,	HEMBB1001905,	HEMBB1001930,	HEMBB1002002,	HEMBB1002009,
	HEMBB1002044,	HEMBB1002255,	HEMBB1002266,	HEMBB1002340,	HEMBB1002458,
20	HEMBB1002492,	HEMBB1002502,	HEMBB1002509,	HEMBB1002510.	HEMBB1002522,
20	HEMBB1002550,	HEMBB1002600, HE	•	BB1002686, HEMBB1	•
	HEMBB1002705,	MAMMA1000175,	MAMMA1000221,	MAMMA1000309,	MAMMA1000424,
	MAMMA1000458,	MAMMA1000468,	MAMMA1000490,	MAMMA1000616,	MAMMA1000621.
0.5	MAMMA1000843,	MAMMA1000845,	MAMMA1000855,	MAMMA1000865,	MAMMA1001030;
25	MAMMA1001075,	MAMMA1001091,	MAMMA1001110,	MAMMA1001222,	MAMMA1001322,
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30	MAMMA1002118,	MAMMA1002156,	MAMMA1002170,	MAMMA1002268,	MAMMA1002269,
	MAMMA1002385,	MAMMA1002566,	MAMMA1002619,	MAMMA1002650,	MAMMA1002685,
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45	NT2RM1000648,			RM1000669, NT2RM1	
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	NT2RM1000811,	NT2RM1000833,	NT2RM1000850,	NT2RM1000852,	NT2RM1000898,
50	NT2RM1000924,	NT2RM1000978,	NT2RM1001003,	NT2RM1001008,	NT2RM1001043,
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55	NT2RM2000522,	NT2RM2000540,	NT2RM2000556,	NT2RM2000566,	NT2RM2000594,
-	NT2RM2000609,	NT2RM2000639,	NT2RM2000669,	NT2RM2000691,	NT2RM2000718,

	NT2RM2000951,	NT2RM2001035,	· NT2RM2001152,	NT2RM2001256,	NT2RM2001291,
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	NT2RM2001547,	NT2RM2001588,	· NT2RM2001592,	NT2RM2001613,	NT2RM2001637,
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10	NT2RM2001982,	NT2RM2001983,	NT2RM2002004,	NT2RM2002014,	NT2RM2002055,
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15	NT2RM4000486,	NT2RM4000514,	NT2RM4000520,	NT2RM4000531,	NT2RM4000532,
70	NT2RM4000534,	NT2RM4000590,	NT2RM4000611,	NT2RM4000717,	NT2RM4000778,
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	NT2RM4001217,	NT2RM4001256,	NT2RM4001258,	NT2RM4001344,	NT2RM4001384,
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25	NT2RM4001938,	NT2RM4001969,	NT2RM4001984,	NT2RM4002018,	NT2RM4002062,
	NT2RM4002073,	NT2RM4002075,	NT2RM4002093,	NT2RM4002128,	NT2RM4002146,
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40	NT2RP1000916,	NT2RP1000943,	NT2RP1000947,	NT2RP1000954,	NT2RP1000980,
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45	NT2RP1001482,	NT2RP1001494,	NT2RP1001616,	NT2RP1001665,	NT2RP2000032,
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	NT2RP2000161,	NT2RP2000239,	NT2RP2000248,	NT2RP2000274,	NT2RP2000288,
	NT2RP2000298,	NT2RP2000310,	NT2RP2000337,	NT2RP2000369,	NT2RP2000420,
50	NT2RP2000422,	NT2RP2000448,	NT2RP2000503,	NT2RP2000510,	NT2RP2000516,
50	NT2RP2000523,	NT2RP2000634,	NT2RP2000656,	NT2RP2000658,	NT2RP2000678,
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E E	NT2RP2001094,	NT2RP2001173,	NT2RP2001196,	NT2RP2001218,	NT2RP2001245,
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	PLACE1010547,	PLACE1010562,	PLACE1010579,	PLACE1010599,	PLACE1010616,
	PLACE1010622,	PLACE1010624,	PLACE1010628,	PLACE1010630,	PLACE1010631,
	PLACE1010662,	PLACE1010714,	PLACE1010739,	PLACE1010743,	PLACE1010800,
55	PLACE1010802,	PLACE1010811,	PLACE1010857,	PLACE1010870,	PLACE1010877,

	PLACE1010891,	PLACE1010896,	PLACE1010916,	PLACE1010917,	PLACE1010960,
	PLACE1010965,	PLACE1011026, PL	ACE1011057, PLAC	E1011143, PLACE1	011160,
_	PLACE1011165,	PLACE1011203,	PLACE1011214,	PLACE1011221,	PLACE1011229,
5	PLACE1011263,	PLACE1011273,	PLACE1011310,	PLACE1011325,	PLACE1011332,
	PLACE1011375,	PLACE1011399,	PLACE1011419,	PLACE1011433,	PLACE1011465,
	PLACE1011503,	PLACE1011520,	PLACE1011563,	PLACE1011635,	PLACE1011641,
10	PLACE1011643,	PLACE1011664,	PLACE1011675,	PLACE1011729,	PLACE1011762,
70	PLACE1011778,	PLACE1011858,	PLACE1011874,	PLACE1011875,	PLACE1011896,
	PLACE1011964,	PLACE1011982,	PLACE1012031,	PLACE2000006,	PLACE2000014,
	PLACE2000015,	PLACE2000021,	PLACE2000033,	PLACE2000061,	PLACE2000072,
15	PLACE2000097,	PLACE2000115,	PLACE2000136,	PLACE2000164,	PLACE2000223,
	PLACE2000371,	PLACE2000379,	PLACE2000477,	PLACE3000059,	PLACE3000121,
	PLACE3000160,	PLACE3000199,	PLACE3000218,	PLACE3000244,	PLACE3000310,
	PLACE3000320,	PLACE3000339,	PLACE3000350,	PLACE3000373,	PLACE4000089,
20	PLACE4000093,	PLACE4000147,	PLACE4000252,	PLACE4000270,	PLACE4000367,
	PLACE4000392,	PLACE4000401,	PLACE4000558,	PLACE4000590,	PLACE4000654,
	PLACE4000670,	SKNMC1000011,	SKNMC1000013,	SKNMC1000046,	SKNMC1000050,
	SKNMC1000091,	THYR01000035,	THYR01000111,	THYRO1000121,	THYR01000124,
25	THYR01000129,	THYR01000156,	THYR01000199,	THYR01000242,	THYR01000270,
	THYR01000279,	THYR01000288,	THYR01000327,	THYR01000381,	THYR01000438,
	THYR01000488,	THYR01000501,	THYR01000502,	THYR01000505,	THYR01000570,
	THYR01000596,	THYRO1000605, TH	YR01000641, THYR	01000662, THYR01	000734,
30	THYR01000777,	THYRO1000783,	THYR01000793,	THYR01000796,	THYR01000805,
	THYR01000829,	THYR01000852,	THYR01000895,	THYR01000934,	THYR01000952,
	THYR01001033,	THYRO1001100,	THYR01001134,	THYRO1001142,	THYR01001204,
	THYR01001262,	THYRO1001271,	THYR01001290,	THYR01001347,	THYR01001559,
35	THYRO1001661,	THYR01001721,	THYR01001745,	Y79AA1000328,	Y79AA1000420,
	Y79AA1000705,	Y79AA1000734,	Y79AA1000748,	Y79AA1000752,	Y79AA1000774,
	Y79AA1000784,	Y79AA1000802,	Y79AA1000824,	Y79AA1000827,	Y79AA1000976,
	Y79AA1001078,	Y79AA1001281,	Y79AA1001312,	Y79AA1001493,	Y79AA1001541,
40	Y79AA1001581,	Y79AA1001585,	Y79AA1001696,	Y79AA1001781,	Y79AA1001848,
	Y79AA1001874,	Y79AA1001923,	Y79AA1002027,	Y79AA1002115,	Y79AA1002139,
	Y79AA1002208, Y79AA1002307.	Y79AA1002209,	Y79AA1002210,	Y79AA1002220,	Y79AA1002298,
	Y79AA1002307,	Y79AA1002311,	Y79AA1002407,	Y79AA1002416,	Y79AA1002431,
45	113001002401				

[0204] Genes exhibiting characteristic features in the expression thereof were selected by statistical analysis of these data. Two examples are shown below to describe the selection of genes of which expression is varied greatly among tissues. The β' actin gene is used frequently as a control in gene expression analysis. Genes of which expression is varied greatly among tissues as compared that of the β-actin gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of β-actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance S_a ². Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance S_b ². By taking variance ratio F as $F = S_b$ ²/ S_a ², genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below by the corresponding clone names: HEMBA1002113, HEMBA1005296, HEMBA1007121, HEMBB1000637, HEMBB1000915, MAMMA1000597, MAMMA1000605, MAMMA1000962, MAMMA1001139, MAMMA1001198, MAMMA1002858, NT2RM2001896, NT2RP2002710, NT2RP2004339, NT2RP2004538, NT2RP3000348, NT2RP3003121, PLACE3000009,

PLACE3000254, THYRO1000569, Y79AA1000131.

[0205] Gene of OVARC1000037 {heterogeneous nuclear ribonucleoprotein (hnRNP)} which expression is varied little. Genes of which expression is varied greatly among tissues as compared that of the OVARC1000037 gene were determined as follows. Specifically, sum of squared deviation was calculated in the signal intensity of β -actin observed in each tissue, which was divided by 7 degrees of freedom to determine a variance S_a^2 . Next, sum of squared deviation was calculated in the signal intensity of a gene to be compared observed in each tissue, which was divided by 7 degrees of freedom to determine a variance S_b^2 . By taking variance ratio F as $F = S_b^2 / S_a^2$, genes with a significance level of 5% or more were extracted in the F distribution. Genes extracted are indicated below, by the corresponding clone names:

4.0					
10	HEMBA1000304,	NT2RM2001716,	NT2RM2001840,	HEMBA1001051,	HEMBA1001109,
	OVARC1001731,	HEMBA1000726,	HEMBA1001286,	HEMBA1000387,	HEMBA1000519,
	NT2RM2001896,	HEMBA1000042,	HEMBA1001085,	HEMBA1001330,	OVARC1000576_
	HEMBA1000575,	NT2RM2000599,	NT2RM2000714,	HEMBA1000469,	NT2RM4000366,
15	HEMBA1001377,	HEMBA1000769,	HEMBA1000338,	NT2RM2000795,	HEMBA1001299,
	HEMBA1000508,	HEMBA1000150,	HEMBA1000774,	HEMBA1001226,	HEMBA1000960:
	NT2RM4000795,	HEMBA1002162,	NT2RM4001876,	NT2RM4002482,	HEMBA1001678,
	HEMBA1002113,	NT2RM4002383,	HEMBA1002229,	HEMBA1002818,	HEMBA1001454,
20	NT2RM4000764,	HEMBA1001510,	HEMBA1001714,	HEMBA1002150,	NT2RM4002044,
	HEMBA1002728,	NT2RM4002189,	HEMBA1001991,	HEMBA1002166,	NT2RM4002499,
	NT2RM4001140,	NT2RM4002504,	HEMBA1002590,	HEMBA1001435,	PLACE1000706,
0.5	HEMBA1002160,	HEMBA1001824,	HEMBA1001463,	HEMBA1001533,	HEMBA1001570,
25	PLACE1001036,	HEMBA1001651,	HEMBA1002381,	HEMBA1002934,	HEMBA1003370,
	HEMBA1003021,	HEMBA1003166,	NT2RP1000738,	NT2RP2000040,	HEMBA1004164,
	HEMBA1003836,	HEMBA1004267,	NT2RP2000845,	HEMBA1003041,	HEMBA1003571,
20	HEMBA1003758,	NT2RP2000108,	HEMBA1003838,	NT2RP1000357,	HEMBA1003376,
30	PLACE1003528,	HEMBA1003528,	NT2RP1001475,	HEMBA1004049,	HEMBA1003212,
	HEMBA1003667,	PLACE1004149,	HEMBA1003926,	HEMBA1004306,	HEMBA1004024,
	NT2RP1000363,	HEMBA1003033,	HEMBA1004335,	HEMBA1003348,	HEMBA1003034,
05	NT2RP2001081,	HEMBA1004056, HE	MBA1003314, HEMB	A1003827, HEMBA1	
35	NT2RP2001036,	NT2RP2001168,	NT2RP2001328,	HEMBA1005035,	NT2RP2001569,
	NT2RP2002439,	HEMBA1005511,	HEMBA1005999,	NT2RP2002862,	NT2RP2002979,
	NT2RP2001394,	HEMBA1004753,	NT2RP2002621,	HEMBA1005853,	HEMBA1005443,
40	NT2RP2002980,	NT2RP2001347,	HEMBA1005241,	NT2RP2002750,	NT2RP2003533,
40	HEMBA1005634,	NT2RP2003034,	HEMBA1006138,	NT2RP2003117,	NT2RP2001366,
	HEMBA1005079,	NT2RP2003293,	NT2RP2002710,	HEMBA1005911,	NT2RP2002752,
	HEMBA1006036,	NT2RP2002987,	HEMBA1006100,	HEMBA1004460,	HEMBA1004538,
45	NT2RP2001943,	NT2RP2002033,	HEMBA1005296,	HEMBA1005829,	HEMBA1005520,
45	HEMBA1005123,	HEMBA1005552,	HEMBA1004930,	NT2RP2001312,	HEMBA1005304,
	HEMBA1005834,	HEMBA1005990,	HEMBA1005526,	NT2RP2003073,	HEMBA1005331,
	HEMBA1006744,	HEMBA1006780,	NT2RP2004339,	HEMBB1000173,	HEMBA1007113,
50	NT2RP2005908,	HEMBB1000376,	HEMBB1000024,	HEMBB1000510,	NT2RP2004580,
50					,

	NT2RP2003912,	HEMBA1006381,	NT2RP2004270,	HEMBA1006993,	NT2RP2005325,
	HEMBB1000218,	HEMBA1007121,	HEMBB1000434,	NT2RP2006166,	PLACE1010310,
	HEMBB1000589,	NT2RP2003986,	HEMBA1006419,	HEMBB1000141,	HEMBB1000030,
5	HEMBB1000099,	HEMBA1006624,	NT2RP2003988,	NT2RP2005753,	HEMBA1007147.
	HEMBB1000354,	NT2RP2006023,	HEMBB1000441,	HEMBB1000523,	HEMBB1000059.
	HEMBA1006631,	NT2RP2004538,	HEMBA1006695,	HEMBB1000490,	NT2RP2006184.
	HEMBB1000573,	NT2RP2004675,	HEMBA1007078,	NT2RP2005457,	NT2RP2005491.
10	HEMBB1000491,	HEMBB1000575, PLA	CE1008405, NT2R	RP2005773, NT2RP2	- · · ·
	HEMBB1000337,	HEMBB1000008,	NT2RP2005701,	NT2RP2003825,	HEMBB1000215,
	HEMBB1000258,	NT2RP2005496,	HEMBB1000374,	NT2RP2005581,	HEMBB1000018,
	HEMBB1000493,	HEMBB1000554,	HEMBB1000671,	NT2RP3000348,	HEMBB1001267,
15	HEMBB1002092,	HEMBB1001749,	PLACE3000304,	HEMBB1001834,	HEMBB1001142.
	HEMBB1001177,	HEMBB1000840,	HEMBB1001436,	HEMBB1001921,	HEMBB1002005,
	NT2RP3001398,	HEMBB1002069,	HEMBB1002094,	HEMBB1001836,	HEMBB1001114,
	NT2RP3000628,	HEMBB1001367,	HEMBB1001588,	HEMBB1002249,	HEMBB1000684,
20	NT2RP2006571,	HEMBB1001802,	HEMBB1000915,	HEMBB1000996,	NT2RP3001274,
20	HEMBB1001564,	NT2RP3000968,	PLACE3000156,	HEMBB1002045,	NT2RP3000341,
	HEMBB1000917,	PLACE1011090,	NT2RP3000742,	PLACE2000435,	HEMBB1001944,
	PLACE3000254,	HEMBB1002218,	NT2RP2006436,	NT2RP3000320,	HEMBB1001126,
	HEMBB1000749,	NT2RP3000644,	HEMBB1001527,	NT2RP3000850,	HEMBB1001967;
25	PLACE3000401,	HEMBB1000637,	HEMBB1001234,	NT2RP3000080,	PLACE2000411,
	PLACE3000009,	PLACE3000070,	HEMBB1002134,	PLACE3000475,	PLACE4000192,
	HEMBB1002520,	HEMBB1002545,	MAMMA1000155,	MAMMA1000307,	THYR01000787,
	MAMMA1000348,	MAMMA1000372,	NT2RP3003121,	MAMMA1000851,	MAMMA1000501,
30	THYR01000569,	MAMMA1000576,	MAMMA1000605,	HEMBB1002442,	HEMBB1002596,
	MAMMA1000198,	MAMMA1000714,	MAMMA1000356,	MAMMA1000760,	MAMMA1000416,
	MAMMA1000931,	MAMMA1000962,	HEMBB1002387,	NT2RP3001712,	HEMBB1002556,
	HEMBB1002617,	MAMMA1000171, PLA	CE4000612, MAMM	MAMMA1 (MAMMA)	000421,
35	MAMMA1000444,	HEMBB1002358,	HEMBB1002453,	HEMBB1002664,	HEMBB1002699,
	MAMMA1000720,	MAMMA1000360,	MAMMA1000744,	MAMMA1000802,	MAMMA1000839,
	MAMMA1000478,	MAMMA1000877,	MAMMA1000585,	MAMMA1000941,	NT2RP3002909,
	MAMMA1000968,	HEMBB1002457,	HEMBB1002495,	MAMMA1000057,	MAMMA1000696,
40	MAMMA1000942,	MAMMA1000302,	MAMMA1000943,	MAMMA1000998,	NT2RP3001752,
	NT2RP3003032,	MAMMA1000429,	MAMMA1000565,	MAMMA1000594,	MAMMA1000625,
	NT2RP3002146,	MAMMA1000084,	MAMMA1000257,	NT2RP3003157,	MAMMA1000431,
	MAMMA1000597,	MAMMA1001078,	MAMMA1001126,	THYR01001537,	NT2RP3004093,
45	MAMMA1001202,	NT2RP3003842,	Y79AA1000342,	MAMMA1002311,	MAMMA1001969,
,,,	Y79AA1000560,	MAMMA1002359,	MAMMA1002056,	MAMMA1002413,	MAMMA1002454,
	MAMMA1002612,	MAMMA1001411,	MAMMA1001035,	MAMMA1001080,	MAMMA1001133,
	NT2RP3004095, MAMMA1001745,	MAMMA1001161,	MAMMA1001203,	MAMMA1001663,	NT2RP3003819,
50	NT2RP4000321.	MAMMA1001790,	NT2RP3004470,	NT2RP3004503,	MAMMA1002293,
50	NT2RP4000515,	MAMMA1001880, MAMMA1002215,	Y79AA1000410,	MAMMA1001970, NT2RP3003576	MAMMA1002058,
	MAMMA1001576,	Y79AA1000131,	MAMMA1002617, MAMMA1001330.	NT2RP3003576,	MAMMA1001139,
	MAMMA1001576,	MAMMA1002125,	MAMMA1001330, MAMMA1002174,	MAMMA1001769, NT2PP4000614	MAMMA1001383,
	NT2RP3004110,	MAMMA1002125, MAMMA1001271,	MAMMA1002174,	NT2RP4000614,	NT2RP4000243,
55	1112IG 2004110'	mumutuvituti,	mammatouleso,	MAMMA1001992,	MAMMA1002033,

	MAMMA1002428,	MAMMA1002590,	MAMMA1001186,	MAMMA1002267,	MAMMA1002322,
	MAMMA1001956,	MAMMA1002155, NT	2RP4000210, MAMM	A1002622, NT2RP3	004125,
_	MAMMA1001220,	MAMMA1001683,	NT2RP3004348,	Y79AA1000214,	Y79AA1000833,
5	NT2RP4000212,	MAMMA1002230,	MAMMA1001452,	MAMMA1001620,	MAMMA1001256,
	MAMMA1001760,	NT2RP3004349,	MAMMA1001783,	MAMMA1001907,	MAMMA1002009,
	MAMMA1002545,	NT2RP4000214,	NT2RP4000728,	MAMMA1001465,	MAMMA1001154,
	MAMMA1001198,	MAMMA1001343,	MAMMA1002310,	NT2RP4000035,	NT2RP4000833,
10	MAMMA1003150,	MAMMA1002886,	NT2RP4001938,	NT2RM2000260,	MAMMA1002629,
	MAMMA1002973,	MAMMA1002721,	MAMMA1002909,	NT2RP4001100,	NT2RM1000857,
	NT2RP4000878,	MAMMA1002844,	NT2RM1000039,	NT2RP4001174,	MAMMA1002665,
	MAMMA1003047,	NT2RM1000086,	NT2RM1000260,	NT2RM1000355,	MAMMA1002701,
15	NT2RP4000918,	MAMMA1002830,	MAMMA1002970,	NT2RP4001677,	NT2RM2000422,
	MAMMA1003004,	MAMMA1002673,	MAMMA1003031,	MAMMA1002764,	MAMMA1002858,
	NT2RP4001679,	NT2RP4002888,,	MAMMA1002711,	NT2RP4001276,	NT2RM1000018,
	NT2RP4001568,	NT2RM1000883.			
20					

[0206] Thus, characteristic features in the expression of a gene are illustrated by comparing and _ statistically analyzing the expression of many genes.

Analysis of disease-associated genes

[0207] Non-enzymic protein glycation reaction is believed to be a cause of a variety of chronic diabetic complications. Accordingly, genes of which expression is elevated or decreased in a glycated protein-specific manner in the endothelial cells are associated with diabetic complications caused by glycated proteins. Vascular endothelial cells are affected with glycated proteins present in blood. Reaction products of non-enzymic protein glycation include amadori compound (glycated protein) as a mildly glycated protein and advanced glycation endproduct as a heavily glycated protein. Hence, a survey was carried out for genes of which expression levels are varied depending on the presence of these glycated proteins in endothelial cells. The mRNAs were extracted from endothelial cells that were cultured in the presence or absence of glycated protein. The mRNAs were converted into radiolabeled first strand cDNAs for preparing probes. The probes were hybridized to the above-mentioned DNA array. Signal of each DNA spot was detected by BAS2000 and analyzed by ArrayGauge (Fuji Photo Film Co., Ltd.).

[0208] Advanced glycation endproduct of bovine serum albumin was prepared as follows: bovine serum albumin (BSA; Sigma) was incubated in a phosphate buffer solution containing 50 mM glucose at 37°C for 8 weeks; and the resulting brownish BSA was dialyzed against a phosphate buffer solution.

[0209] Human normal pulmonary arterial endothelial cells (Cell Applications) were cultured in an Endothelial Cell Growth Medium (Cell Applications). The culture dish (Farcon) with the cells were incubated in a CO₂ incubator (37°C, 5% CO₂, in a humid atmosphere). When the cells were grown to be confluent in the dish, 250 µg/ml of bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin was added thereto and the cells were incubated for 33 hours. The mRNA was extracted from the cells by using a FastTack^(TM) 2.0 kit (Invitrogen). The labeling of hybridization probe was carried out by using the mRNA according to the same procedure as described above.

[0210] Table 169 shows the expression level of each cDNA in human pulmonary arterial endothelial cells cultured in a medium containing bovine serum albumin (sigma), glycated bovine serum albumin (Sigma) or advanced glycation endproduct of bovine serum albumin. Genes of which expression was detected in the endothelial cell are as follows:

55

50

	HEMBA1000005.	LIEMPA1000010	UTM # 100000	ITTIO 4 1 0 0 0 0 4 0	ITIMA 10000 to
		HEMBA1000012,	HEMBA1000020,	HEMBA1000042,	HEMBA1000046,
	HEMBA1000076,	HEMBA1000111,	HEMBA1000150,	HEMBA1000185,	HEMBA1000216,
_	HEMBA1000231,	HEMBA1000280,	HEMBA1000282,	HEMBA1000288,	HEMBA1000304,
5	HEMBA1000327,	HEMBA1000338,	HEMBA1000356,	HEMBA1000357,	HEMBA1000366,
	HEMBA1000376,	HEMBA1000387,	HEMBA1000428,	HEMBA1000456,	HEMBA1000459,
	HEMBA1000460,	HEMBA1000469,	HEMBA1000501,	HEMBA1000504,	HEMBA1000505,
	HEMBA1000508,	HEMBA1000519,	HEMBA1000534,	HEMBA1000540,	HEMBA1000542,
10	HEMBA1000555,	HEMBA1000557,	HEMBA1000575,	HEMBA1000592,	HEMBA1000604,
	HEMBA1000622,	HEMBA1000636,	HEMBA1000655,	HEMBA1000673,	HEMBA1000682,
	HEMBA1000686,	HEMBA1000702,	HEMBA1000726,	HEMBA1000727,	HEMBA1000749,
	HEMBA1000769,	HEMBA1000774,	HEMBA1000791,	HEMBA1000827,	HEMBA1000852,
15	HEMBA1000872,	HEMBA1000876,	HEMBA1000918,	HEMBA1000942,	HEMBA1000960,
	HEMBA1000974,	HEMBA1001020,	HEMBA1001051,	HEMBA1001059,	HEMBA1001060,
	HEMBA1001080,	HEMBA1001085,	HEMBA1001088,	HEMBA1001109,	HEMBA1001122,
	HEMBA1001123,	HEMBA1001140,	HEMBA1001172,	HEMBA1001196,	HEMBA1001197,
	HEMBA1001213,	HEMBA1001226,	HEMBA1001235,	HEMBA1001247,	HEMBA1001265,
20	HEMBA1001281,	HEMBA1001286,	HEMBA1001299,	HEMBA1001302,	HEMBA1001326,
	HEMBA1001330,	HEMBA1001351,	HEMBA1001377,	HEMBA1001387,	HEMBA1001398,
	HEMBA1001432,	HEMBA1001433,	HEMBA1001435,	HEMBA1001446,	HEMBA1001450,
	HEMBA1001454,	HEMBA1001463, HE	MBA1001497, HEME	8A1001510, HEMBA1	
25	HEMBA1001526,	HEMBA1001533,	HEMBA1001557,	HEMBA1001566,	HEMBA1001569,
	HEMBA1001570,	HEMBA1001579,	HEMBA1001581,	HEMBA1001595,	HEMBA1001608,
	HEMBA1001620,	HEMBA1001640,	HEMBA1001647,	HEMBA1001651,	HEMBA1001655,
	HEMBA1001658,	HEMBA1001678,	HEMBA1001714,	HEMBA1001718,	HEMBA1001723,
30	HEMBA1001746,	HEMBA1001784,	HEMBA1001791,	HEMBA1001800,	HEMBA1001804,
	HEMBA1001809,	HEMBA1001819,	HEMBA1001824,	HEMBA1001844,	HEMBA1001866,
	HEMBA1001888,	HEMBA1001912,	HEMBA1001913,	HEMBA1001918,	HEMBA1001940,
	HEMBA1001960,	HEMBA1001987,	HEMBA1001991,	HEMBA1002003,	HEMBA1002008,
25	HEMBA1002049,	HEMBA1002100,	HEMBA1002102,	HEMBA1002113,	HEMBA1002119,
35	HEMBA1002125,	HEMBA1002151,	HEMBA1002153,	HEMBA1002160,	HEMBA1002161,
	HEMBA1002166,	HEMBA1002185,	HEMBA1002189,	HEMBA1002191,	HEMBA1002226,
	HEMBA1002229,	HEMBA1002237,	HEMBA1002241,	HEMBA1002267,	HEMBA1002270,
	HEMBA1002337,	HEMBA1002348,	HEMBA1002381,	HEMBA1002458,	HEMBA1002462,
40	HEMBA1002469,	HEMBA1002503,	HEMBA1002508,	HEMBA1002542,	HEMBA1002547,
	HEMBA1002552,	HEMBA1002558,	HEMBA1002583,	HEMBA1002590,	HEMBA1002592,
	HEMBA1002609,	HEMBA1002624,	HEMBA1002629,	HEMBA1002645,	HEMBA1002678,
	HEMBA1002688,	HEMBA1002703,	HEMBA1002712,	HEMBA1002728,	HEMBA1002730,
45	HEMBA1002750,	HEMBA1002770,	HEMBA1002779,	HEMBA1002780,	HEMBA1002818,
	HEMBA1002833,	HEMBA1002876,	HEMBA1002934,	HEMBA1002935,	HEMBA1002968,
	HEMBA1002970,	HEMBA1002973, HE	MBA1003021, HEMB	A1003033, HEMBA1	003034,
	HEMBA1003041,	HEMBA1003046,	HEMBA1003064,	HEMBA1003067,	HEMBA1003079,
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50	HEMBA1003142,	HEMBA1003166,	HEMBA1003175,	HEMBA1003179,	HEMBA1003202,
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	Y79AA1000774,	Y79AA1000782,	Y79AA1000784,	Y79AA1000827,	Y79AA1000833,

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5	Y79AA1001211,	Y79AA1001216,	Y79AA1001228,	Y79AA1001236,	Y79AA1001299,
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10	Y79AA1001846,	Y79AA1001866,	Y79AA1001875,	Y79AA1001923,	Y79AA1001963,
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	Y79AA1002351,	Y79AA1002361, Y79	9AA1002472, Y79A	A1002482	·

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[0211] Signal ratios of EC_AGE_BSA to EC_BSA and of EC_glycated_BSA to EC_BSA were calculated for each gene. Genes with high signal ratios were selected. In the case of calculating the ratio of signal value of 40 or less to that of more than 40, such signal values were, for convenience, taken as 40 instead of the real values. When the ratio EC_AGE_BSA/EC_BSA is 2 or more, expression of the genes exhibiting such ratio is expected to be elevated due to advanced glycation endproduct of bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC_AGE_BSA/EC_BSA ranges from 0.5 to 2, expression of the genes exhibiting such ratio is expected to be unaffected due to advanced glycation endproduct of bovine serum albumin. When the ratio EC_AGE_BSA/EC_BSA is less than 0.5, expression of the genes exhibiting such ratio value is expected to be decreased due to advanced glycation endproduct of bovine serum albumin. The lower the value is, the lower the gene expression level is. [0212] Clone with EC_AGE_BSA/EC_BSA ratio of 2 or higher are as follows: HEMBA1003958, MAMMA1001256, PLACE2000411.

[0213] Clone with EC_AGE_BSA/EC_BSA ratio of 0.5 or less is as follows: MAMMA1001783. These cDNAs are associated with diabetes.

[0214] When the ratio EC_glycated_BSA/EC_BSA is 2 or more, the expression level of the gene exhibiting such ratio is expected to be elevated due to glycated bovine serum albumin. The higher the value is, the higher the gene expression level is. When the ratio EC_glycated_BSA/EC_BSA ranges from 0.5 to 2, the expression level of the gene exhibiting such ratio is expected to be unaffected with glycated bovine serum albumin. When the ratio EC_glycated_BSA/EC_BSA is less than 0.5, the expression level of a gene exhibiting such ratio is expected to be decreased due to glycated bovine serum albumin. The lower the value is, the lower the gene expression level is.

[0215] Clones with EC_glycated_BSA/EC_BSA ratio of 2 or more are as follows: HEMBA1004850, MAMMA1001256, MAMMA1002132 and PLACE3000119.

[0216] A clone with EC_glycated_BSA/EC_BSA ratio of 0.5 or less is as follows: MAMMA1001783.

[0217] These cDNAs are also associated with diabetes.

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Analysis of genes associated with neural cell differentiation

[0218] Genes involved in neural cell differentiation are useful for treating neurological diseases. It is possible that genes with varying expression levels in response to induction of cellular differentiation in neural cells are associated with neurological diseases.

[0219] A survey was performed for genes of which expression levels are varied in response to induction of differentiation (stimulation by retinoic acid (RA)) in cultured cells of a neural strain, NT2.

[0220] The NT2 cells were treated basically according to supplier's instruction manual. "Undifferentiated NT2 cells" means NT2 cells successively cultured in an Opti-MEM I (GIBCO-BRL; catalog No. 31985) containing 10%(v/v) fetal bovine serum and 1%(v/v) penicillin-streptomycin (GIBCO BRL). "NT2 cells cultured in the presence of retinoic acid" means the cells resulted from transferring undifferentiated NT2 cells into a retinoic acid-containing medium, which consists of D-MEM (GIBCO BRL; catalog No. 11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin and $10~\mu$ M retinoic acid (GIBCO-BRL), and the subsequent successive culture therein for 5 weeks. "NT2 cells that were cultured in the presence of retinoic acid and then further cultured in the presence of cell-division inhibitor added" means NT2 cells resulted from transferring NT2 cells cultured in the presence of retinoic acid for 5 weeks into a cell-division inhibitor-containing medium, which consisted of D-MEM(GIBCO BRL; catalog No.11965), 10%(v/v) fetal bovine serum, 1%(v/v) penicillin-streptomycin, 10μ M retinoic acid, $10~\mu$ M FudR (5-fluoro-2'-deoxyuridine: GIBCO BRL), $10~\mu$ M Urd (Uridine: GIBCO BRL) and $1~\mu$ M araC (Cytosine β-D-Arabinofuranoside: GIBCO BRL), and the subsequence succes-

sive culture for 2 weeks. Each of the cells were treated with trypsin and then harvested. Total RNAs were extracted from the cells by using S.N.A.P.(TM) Total RNA Isolation kit (Invitrogen^(r)). The labeling of probe used for hybridization was carried out by using 10 µg of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of stimulation for inducing differentiation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment-of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis, clones with the difference can be statistically detected even when the signals were low. Accordingly, clones with signal value of 40 or less were also assessed for the selection. [0221] Tables 170-349 show the expression level of each cDNA in undifferentiated NT2 cells, NT2 cells cultured in the presence of RA, and NT2 cells that were cultured in the presence of cell-division inhibitor added.

[0222] Averaged signal values (M_1, M_2) and sample variances (s_1^2, s_2^2) were calculated for each gene in each of the cells, and then, the pooled sample variances s^2 were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula: $t=(M_1-M_2)/s/(1/3+1/3)^{1/2}$. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information on an increase (+) or decrease (-) in the expression level of a gene in the treated cells when the level is compared with that of untreated undifferentiated cells.

20 [0223] Clones of which expression levels increased by RA are as follows: HEMBA1000005, HEMBA1000042, HEMBA1000046, HEMBA1000076. HEMBA1000111, HEMBA1000141, HEMBA1000150, HEMBA1000185. HEMBA1000282. HEMBA1000304. HEMBA1000307. HEMBA1000338, HEMBA1000357. HEMBA1000376. HEMBA1000387, HEMBA1000392, HEMBA1000428, HEMBA1000456, HEMBA1000459. HEMBA1000469, HEMBA1000504. HEMBA1000508, HEMBA1000519, HEMBA1000540, HEMBA1000545. HEMBA1000557. HEMBA1000563, HEMBA1000568, HEMBA1000575, HEMBA1000588, HEMBA1000592, HEMBA1000604, HEMBA1000622, HEMBA1000655, HEMBA1000673, HEMBA1000682, HEMBA1000726. HEMBA1000727, HEMBA1000749, HEMBA1000769, HEMBA1000774, HEMBA1000791, HEMBA1000822, HEMBA1000872. HEMBA1000876, HEMBA1000910, HEMBA1000942, HEMBA1000943. HEMBA1000960, HEMBA1000972, HEMBA1000974, HEMBA1000991, HEMBA1001008. HEMBA1001020, 30 HEMBA1001043, HEMBA1001051, HEMBA1001060, HEMBA1001071. HEMBA1001077, HEMBA1001085, HEMBA1001094, HEMBA1001109, HEMBA1001121. HEMBA1001122, HEMBA1001140, HEMBA1001172. HEMBA1001226, HEMBA1001235, HEMBA1001265, HEMBA1001281, HEMBA1001294, HEMBA1001299. HEMBA1001319, HEMBA1001323, HEMBA1001330, HEMBA1001351, HEMBA1001361, HEMBA1001377. HEMBA1001388, HEMBA1001391, HEMBA1001398, HEMBA1001432, HEMBA1001435, HEMBA1001442, HEMBA1001454, HEMBA1001455, HEMBA1001497, HEMBA1001517, HEMBA1001569, HEMBA1001570, HEMBA1001581, HEMBA1001585, HEMBA1001620, HEMBA1001711, HEMBA1001718, HEMBA1001723, HEMBA1001761, HEMBA1001815, HEMBA1001819, HEMBA1001861, HEMBA1001864, HEMBA1001869, HEMBA1001888, HEMBA1001915, HEMBA1001918. HEMBA1001940, HEMBA1001964, HEMBA1001967, HEMBA1001979, HEMBA1001987. HEMBA1001991, HEMBA1002008, HEMBA1002022, HEMBA1002039, HEMBA1002102, HEMBA1002049. HEMBA1002084, HEMBA1002113, HEMBA1002144, HEMBA1002160, HEMBA1002162, HEMBA1002185, HEMBA1002212, HEMBA1002226, HEMBA1002229, HEMBA1002267, HEMBA1002270, HEMBA1002337, HEMBA1002381, HEMBA1002458, HEMBA1002477, HEMBA1002508, HEMBA1002558, HEMBA1002561, HEMBA1002583, HEMBA1002590, HEMBA1002628, HEMBA1002645, HEMBA1002661, HEMBA1002678, HEMBA1002712, HEMBA1002728, HEMBA1002780, HEMBA1002850, HEMBA1002886, HEMBA1002934, HEMBA1002935, HEMBA1002939, HEMBA1002951, HEMBA1002968, HEMBA1002970, HEMBA1002973, HEMBA1002999, HEMBA1003021, HEMBA1003033, HEMBA1003034, HEMBA1003064, HEMBA1003067, HEMBA1003078, HEMBA1003086, HEMBA1003096, HEMBA1003129. HEMBA1003142, HEMBA1003148, HEMBA1003166, HEMBA1003175, HEMBA1003197. HEMBA1003199. HEMBA1003202. HEMBA1003204, HEMBA1003212, HEMBA1003235. HEMBA1003250. HEMBA1003273. HEMBA1003276, HEMBA1003278, HEMBA1003291. HEMBA1003309. HEMBA1003322. HEMBA1003328, HEMBA1003348, HEMBA1003369. HEMBA1003376. HEMBA1003384, HEMBA1003395. HEMBA1003463. HEMBA1003480, HEMBA1003595, HEMBA1003531, HEMBA1003548, HEMBA1003591. HEMBA1003597, HEMBA1003617, HEMBA1003621, HEMBA1003622, HEMBA1003637, HEMBA1003640, HEMBA1003645, HEMBA1003646, HEMBA1003656, HEMBA1003679, HEMBA1003692. HEMBA1003715, HEMBA1003720, HEMBA1003725, HEMBA1003729, HEMBA1003758, HEMBA1003803, HEMBA1003805, HEMBA1003836, HEMBA1003838, HEMBA1003879, HEMBA1003885, HEMBA1003893, HEMBA1003908, HEMBA1003937, HEMBA1003942, HEMBA1003953, HEMBA1003958, HEMBA1003959, HEMBA1003978, HEMBA1003987, HEMBA1003989, HEMBA1004000, HEMBA1004011, HEMBA1004012, HEMBA1004015, HEMBA1004024,

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     Y79AA1002258.
     [0226] Clones of which expression levels decrease by RA/inhibitor are as follows:
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                                   HEMBA1006467, HEMBA1006486,
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                                   HEMBB1001665, MAMMA1000684,
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    Y79AA1000346, Y79AA1001541.
    [0227] Clones of which expression levels increase in the presence of both RA and RA/inhibitor are as follows:
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    MAMMA1001243, MAMMA1002769, MAMMA1002871, MAMMA1002941,
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                                                                    THYRO1000934,
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                                                     THYRO1001290,
                                                                     Y79AA1000800.
                                                                                    Y79AA1000976,
     Y79AA1001585, Y79AA1001696, Y79AA1001711, Y79AA1002258.
     [0228] Clones of which expression levels decrease in the presence of both RA and RA/inhibitor are as follows:
    HEMBA1000946, HEMBA1003569, HEMBA1005570, NT2RM1000666.
                                                                    NT2RM2000092, NT2RM2000594,
    NT2RM4001754,
                     NT2RM4001905,
                                     NT2RP2002047,
                                                    NT2RP3000980.
                                                                     NT2RP3002081.
                                                                                    NT2RP3004594,
    NT2RP4001950.
                    NT2RP4002408,
                                    OVARC1000431, OVARC1001942,
                                                                    OVARC1001943,
                                                                                    PLACE1003190.
     PLACE1005923, PLACE1007257, PLACE1010624, Y79AA1000346.
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     [0229] These are neurological disease-associated clones.
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Analysis of rheumatoid arthritis-associated genes

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[0230] The onset of rheumatoid arthritis is thought to be involved in the proliferation of synovial cells covering inner surfaces of joint cavity and in inflammatory reaction resulted from the action of cytokines produced by leukocytes infiltrating into the joint synovial tissues (Rheumatism Information Center, http://www.rheuma-net.or.jp/). Recent studies have also revealed that tissue necrosis factor (TNF)-α participates in the onset (Current opinion in immunology 1999, 11, 657-662). When the expression of a gene exhibits responsiveness to the action of TNF on synovial cells, the gene is considered to be involved in rheumatoid arthritis.

[0231] A survey was performed for genes of which expression levels are varied in response to TNF- α in the primary cell culture of synovial tissue. The primary cultured cells of the smooth muscle (Cell Applications) were grown to be confluent in a culture dish, and then, human TNF- α (Boehringer-Mannheim) was added at a final concentration of 10 ng/ml thereto. The culture was further continued for 24 hours.

[0232] Total RNA was extracted from the cells by using S.N.A.P.($^{\text{TM}}$) Total RNA Isolation kit (Invitrogen). The labeling of probe used for hybridization was carried out by using 10 μ g of the total RNA according to the same methods as described above. The data were obtained in triplicate (n=3). The data of signal value representing gene expression level in the cells in the presence of TNF stimulation were compared with those in the absence of the stimulation. The comparison was performed by statistical treatment of two-sample t-test. Clones with significant difference in the signal distribution were selected under the condition of p<0.05. In this analysis,

clones with the difference can be statistically detected even when the signals were low.

Accordingly, clones with signal value of 40 or less were also assessed for the selection.

Table 352 shows the expression level of each cDNA in synovial cells cultured in the absence or presence of TNF. **[0233]** Averaged signal values (M_1, M_2) and sample variances (s_1^2, s_2^2) for each gene were calculated in each of the cells, and then, the pooled sample variances s^2 were obtained from the sample variances of the two types of cells to be compared. The t-values were determined according to the following formula: $t=(M_1-M_2)/s/(1/3+1/3)^{1/2}$. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01 in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively.

The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the stimulated cells when the level is compared with that of unstimulated cells.

[0234] Clones of which expression levels are elevated by TNF- α are as follows: HEMBA1000005, HEMBA1000012, HEMBA1000020, HEMBA1000046, HEMBA1000076. HEMBA1000111. HEMBA1000168, HEMBA1000185, HEMBA1000201, HEMBA1000231. HEMBA1000243. HEMBA1000280. HEMBA1000282, HEMBA1000304, HEMBA1000307, HEMBA1000327, HEMBA1000356, HEMBA1000376, HEMBA1000387, HEMBA1000390, HEMBA1000418, HEMBA1000460, HEMBA1000491, HEMBA1000501. HEMBA1000518, HEMBA1000519, HEMBA1000520, HEMBA1000531, HEMBA1000534, HEMBA1000542, HEMBA1000545, HEMBA1000591, HEMBA1000592, HEMBA1000594. HEMBA1000636, HEMBA1000655, HEMBA1000657, HEMBA1000673, HEMBA1000682. HEMBA1000686, HEMBA1000722, HEMBA1000726, HEMBA1000827, HEMBA1000870. HEMBA1000918, HEMBA1000971, HEMBA1000974, HEMBA1000986, HEMBA1001019, HEMBA1001043. HEMBA1001051, HEMBA1001059, HEMBA1001060, HEMBA1001071, HEMBA1001080. HEMBA1001109, HEMBA1001140, HEMBA1001172, HEMBA1001196, HEMBA1001213, HEMBA1001226, HEMBA1001281, HEMBA1001299, HEMBA1001302, HEMBA1001303, HEMBA1001323, HEMBA1001326, HEMBA1001327, HEMBA1001330, HEMBA1001351, HEMBA1001407, HEMBA1001411, HEMBA1001446. HEMBA1001454, HEMBA1001569, HEMBA1001647, HEMBA1001714, HEMBA1001800, HEMBA1001804, HEMBA1001809, HEMBA1001888, HEMBA1001912, HEMBA1001921, HEMBA1001967, HEMBA1002084, HEMBA1002161, HEMBA1002166, HEMBA1002241, HEMBA1002337, HEMBA1002363, HEMBA1002389, HEMBA1002458, HEMBA1002460, HEMBA1002469, HEMBA1002538, HEMBA1002542, HEMBA1002547, HEMBA1002609, HEMBA1002624. HEMBA1002659, HEMBA1002750, HEMBA1002770, HEMBA1002779, HEMBA1002810. HEMBA1002816, HEMBA1002818, HEMBA1002850. HEMBA1002863. HEMBA1003021, HEMBA1003033. HEMBA1003078, HEMBA1003166. HEMBA1003202. HEMBA1003204. HEMBA1003229, HEMBA1003235, HEMBA1003276. HEMBA1003286. HEMBA1003296. HEMBA1003370. HEMBA1003376, HEMBA1003403. HEMBA1003418. HEMBA1003433. HEMBA1003447. HEMBA1003560. HEMBA1003569, HEMBA1003571, HEMBA1003591, HEMBA1003597. HEMBA1003598. HEMBA1003621. HEMBA1003656, HEMBA1003662, HEMBA1003680, HEMBA1003715, HEMBA1003725, HEMBA1003729, HEMBA1003733, HEMBA1003742. HEMBA1003773, HEMBA1003783. HEMBA1003950. HEMBA1004012, HEMBA1004015. HEMBA1004048, HEMBA1004074, HEMBA1004086, HEMBA1004111, HEMBA1004131, HEMBA1004202, HEMBA1004203, HEMBA1004207, HEMBA1004248, HEMBA1004274, HEMBA1004321, HEMBA1004330, HEMBA1004356, HEMBA1004366, HEMBA1004405, HEMBA1004408, HEMBA1004429, HEMBA1004499, HEMBA1004507, HEMBA1004509, HEMBA1004542, HEMBA1004596, HEMBA1004604,

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    NT2RP3002876,
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                                                                  NT2RP3004209,
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    NT2RP4000078,
                   NT2RP4000541,
                                   NT2RP4000588.
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    OVARC1000106.
                   OVARC1000151.
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    OVARC1000605.
                   OVARC1000679,
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OVARC1001038.
                    OVARC1001055.
                                    OVARC1001085.
                                                    OVARC1001129.
                                                                    OVARC1001167,
                                                                                    OVARC1001339.
     OVARC1001425.
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                                    OVARC1001762.
                                                    OVARC1001766.
                                                                    OVARC1001942,
                                                                                    OVARC1002044,
     OVARC1002138.
                     PLACE1000004,
                                     PLACE1000005.
                                                     PLACE1000420,
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     PLACE1000653,
                    PLACE1001168,
                                    PLACE1001311,
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                                                                    PLACE1001920.
                                                                                    PLACE1001983.
    PLACE1002066.
                    PLACE1002072,
                                    PLACE1002140,
                                                    PLACE1002171.
                                                                    PLACE1002319,
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     PLACE1002499,
                    PLACE1002532.
                                    PLACE1002665.
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     PLACE1003605,
                    PLACE1003704,
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                                                    PLACE1003885.
                                                                    PLACE1004405.
                                                                                    PLACE1004629.
     PLACE1004686,
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                                    PLACE1005066.
                                                    PLACE1006077.
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     PLACE1006143,
                    PLACE1006325,
                                    PLACE1006488,
                                                    PLACE1006805,
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     PLACE1007858,
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                                    PLACE1011725,
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                                                                    PLACE2000264,
                                                                                    PLACE2000394,
     PLACE2000419.
                    PLACE3000160,
                                    PLACE3000220,
                                                    PLACE3000254,
                                                                    PLACE3000271,
                                                                                    PLACE3000339,
     PLACE3000341,
                    PLACE3000350,
                                    PLACE3000353,
                                                    PLACE3000401,
                                                                    PLACE4000300,
                                                                                    SKNMC1000091,
     THYRO1000855,
                     THYRO1001559,
                                     Y79AA1000065,
                                                     Y79AA1000202,
                                                                     Y79AA1000214,
                                                                                     Y79AA1000346,
     Y79AA1000784, Y79AA1000833, Y79AA1000968, Y79AA1001555, Y79AA1002220.
     [0235] Clones of which expression levels decrease by TNF-\alpha are as follows:
     HEMBA1002150, HEMBB1000240,
                                   NT2RM2000469, NT2RM2000984, NT2RM2001688,
                                                                                    NT2RM4000290,
     NT2RM4000496,
                    NT2RM4000590,
                                    NT2RM4001047,
                                                    NT2RM4001582,
                                                                    NT2RM4001611,
                                                                                    NT2RM4001650.
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     NT2RM4002075,
                    NT2RM4002128.
                                     NT2RP1000174,
                                                    NT2RP1000243,
                                                                    NT2RP1000581.
                                                                                    NT2RP1000688.
     NT2RP1000767.
                    NT2RP1000825,
                                    NT2RP1001185,
                                                    NT2RP1001286.
                                                                    NT2RP1001432.
                                                                                    NT2RP1001457,
     NT2RP2000001,
                    NT2RP2000248,
                                    NT2RP2000841.
                                                    NT2RP2001813.
                                                                    NT2RP2002137,
                                                                                    NT2RP2002928.
     NT2RP2003517,
                    NT2RP2003559.
                                    NT2RP2003564.
                                                    NT2RP2004933,
                                                                    NT2RP2005038.
                                                                                    NT2RP2006365,
     NT2RP3000072.
                    NT2RP3000320.
                                    NT2RP3000484,
                                                    NT2RP3000980.
                                                                    NT2RP3001111,
                                                                                    NT2RP3001420,
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     NT2RP3001495.
                    NT2RP3002056.
                                    NT2RP3002057.
                                                    NT2RP3002545.
                                                                    NT2RP3002713,
                                                                                    NT2RP3002799,
     NT2RP3002869.
                    NT2RP3002953,
                                    NT2RP3002955.
                                                    NT2RP3003282,
                                                                    NT2RP3003290,
                                                                                    NT2RP3003384,
    NT2RP3003385.
                    NT2RP3003870.
                                    NT2RP3004207,
                                                    NT2RP3004262,
                                                                    NT2RP3004527,
                                                                                    NT2RP4000500,
    NT2RP4000524,
                    NT2RP4000787.
                                    NT2RP4000927,
                                                    NT2RP4000955,
                                                                    NT2RP4000989,
                                                                                    NT2RP4001442.
    NT2RP4001638,
                    NT2RP4001950,
                                    NT2RP4002888.
                                                    NT2RP5003524,
                                                                    OVARC1001270,
                                                                                    PLACE1000246,
30
    PLACE1002816.
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[0236] These are rheumatoid arthritis-associated clones.

Analysis of ultraviolet radiation damage-associated genes

[0237] It is known that ultraviolet rays give considerably adverse influence on the health. In recent years, there have been significant risks of tissue damage by ultraviolet rays because of destruction of the ozone layer. Thus, ultraviolet radiation has been recognized as a risk factor for skin diseases such as skin cancers (United States Environmental Protection Agency: Ozone Depletion Home Page, http://www.epa.gov/ozone/). Genes of which expression levels are varied in skin epidermal cells exposed to ultraviolet rays are considered to be associated with skin damage caused by ultraviolet radiation.

[0238] After primary cultured skin fibroblast cells were irradiated with ultraviolet ray and were cultured, a survey was performed for genes of which expression levels were varied depending on the irradiation of ultraviolet ray. First, after cultured to be confluent, the primary cultured skin fibroblast cells (Cell Applications) were exposed to $10,000 \, \mu J/cm^2$ of 254-nm ultraviolet light.

[0239] Messenger RNAs were, then, extracted by using a FastTrack™ 2.0 mRNA Isolation kit (Invitrogen Co.) from the unexposed cells and from the cells that were exposed to the ultraviolet light and then cultured for 4 or 24 hours. The labeling of the hybridization probe was carried out by using a 1.5 μg of each mRNA in the same manner as described above. The data were obtained in triplicate (n=3). The hybridization signals were compared between the cells exposed to the ultraviolet light and the unexposed cells. The comparison was preformed by statistical treatment with two-sample t-test. Clones with significant differences in the signal distribution were selected under the condition of p<0.05. In this analysis, even when the signal is lower than others, the difference in the signal values can be detected statistically. Accordingly, clones with signal value of 40 or lower were also assessed for selection.

[0240] Tables 353-509 show the expression of each cDNA in skin-derived fibroblast cells exposed and unexposed to ultraviolet light.

[0241] Averaged signal values (M₁, M₂) and sample variances (s₁², s₂²) were calculated for each gene in each of the cells, and then, the pooled sample variances s² were obtained from the sample variances of the two types of cells to be compared. The t values were determined according to the following formula: t=(M₁-M₂)/s/(1/3+1/3)^{1/2}. When the determined t-value was greater than a t-value at P, which means the probability of significance level, of 0.05 or 0.01

in the t-distribution table with 4 degrees of freedom, the difference was judged to be found in the expression level of the gene between the two types of cells at p<0.05 or p<0.01, respectively. The tables also include the information of an increase (+) or decrease (-) in the expression level of a gene in the exposed cells in comparison with that of unexposed cells.

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[0242] The expression levels of the following clones were elevated 4 or 24 hours after the ultraviolet irradiation:
     HEMBA1000542, HEMBA1001808, HEMBA1002177, HEMBA1003314, MAMMA1001874, NT2RM2001100,
                                   NT2RP4000657,
     NT2RP2005732,
                    NT2RP3000592,
                                                   OVARC1000004,
                                                                  OVARC1001092,
                                                                                  OVARC1001342.
     PLACE1002816,
                    NT2RM4001002,
                                    NT2RM4001813,
                                                   NT2RM4002266,
                                                                   NT2RP2001174,
                                                                                  NT2RP2001196.
    NT2RP2005358,
                    NT2RP3000690.
                                   NT2RP3001216,
                                                   NT2RP3003464,
                                                                  PLACE1006382,
                                                                                  THYRO1000070,
     THYRO1001100, Y79AA1000342.
     [0243] The expression levels of the following clones were decreased 4 or 24 hours after the ultraviolet irradiation:
    HEMBA1000005,
                    HEMBA1000150.
                                   HEMBA1000156,
                                                   HEMBA1000158.
                                                                  HEMBA1000168.
                                                                                  HEMBA1000231.
    HEMBA1000304,
                    HEMBA1000307,
                                   HEMBA1000333.
                                                   HEMBA1000366.
                                                                  HEMBA1000369,
                                                                                  HEMBA1000390,
     HEMBA1000396,
                    HEMBA1000418,
                                   HEMBA1000434.
                                                   HEMBA1000464,
                                                                  HEMBA1000469,
                                                                                  HEMBA1000490,
    HEMBA1000504,
                    HEMBA1000505.
                                   HEMBA1000557,
                                                   HEMBA1000657,
                                                                  HEMBA1000673,
                                                                                  HEMBA1000682,
    HEMBA1000686.
                    HEMBA1000727,
                                   HEMBA1000752.
                                                   HEMBA1000851,
                                                                  HEMBA1000852,
                                                                                  HEMBA1000870,
    HEMBA1000872,
                    HEMBA1001085,
                                    HEMBA1001121,
                                                   HEMBA1001133,
                                                                  HEMBA1001235,
                                                                                  HEMBA1001265,
    HEMBA1001281.
                    HEMBA1001289,
                                   HEMBA1001299,
                                                   HEMBA1001303,
                                                                  HEMBA1001310,
                                                                                  HEMBA1001323,
    HEMBA1001595,
                    HEMBA1001620,
                                   HEMBA1001640,
                                                   HEMBA1001678,
                                                                  HEMBA1001712,
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    HEMBA1001950,
                    HEMBA1001987,
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                                                   HEMBA1002321,
                                                                  HEMBA1002341,
                                                                                  HEMBA1002419.
    HEMBA1002679,
                    HEMBA1002728,
                                   HEMBA1002818,
                                                   HEMBA1002935,
                                                                  HEMBA1002999,
                                                                                  HEMBA1003034.
    HEMBA1003071,
                    HEMBA1003098,
                                   HEMBA1003142,
                                                   HEMBA1003175,
                                                                  HEMBA1003202.
                                                                                  HEMBA1003212.
    HEMBA1003220,
                    HEMBA1003276.
                                   HEMBA1003373,
                                                   HEMBA1003417.
                                                                  HEMBA1003447,
                                                                                  HEMBA1003528.
    HEMBA1003684,
                    HEMBA1003799,
                                   HEMBA1003885,
                                                   HEMBA1003989.
                                                                  HEMBA1004011,
                                                                                  HEMBA1004055.
25
    HEMBA1004133.
                    HEMBA1004225,
                                   HEMBA1004272.
                                                   HEMBA1004353.
                                                                  HEMBA1004631.
                                                                                  HEMBA1004669.
    HEMBA1004705,
                    HEMBA1004753,
                                                                  HEMBA1004816,
                                   HEMBA1004776.
                                                   HEMBA1004803,
                                                                                  HEMBA1004900,
    HEMBA1005047,
                    HEMBA1005079.
                                   HEMBA1005101.
                                                   HEMBA1005149.
                                                                  HEMBA1005152,
                                                                                  HEMBA1005202.
    HEMBA1005314,
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                    HEMBA1005372,
                                                   HEMBA1005513.
                                                                  HEMBA1005518,
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                                                                  HEMBA1005632.
                                                                                  HEMBA1005853,
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    HEMBA1006031,
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                                   HEMBA1006485,
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    HEMBA1006789.
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    HEMBB1000317,
                    HEMBB1000336,
                                   HEMBB1000354,
                                                   HEMBB1000369,
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    HEMBB1000438,
                    HEMBB1000592,
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    HEMBB1000887,
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                                                                                  HEMBB1002556.
    HEMBB1002600, HEMBB1002664,
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	PLACE1011964,	PLACE1012031,	PLACE2000015,	PLACE2000021,	PLACE2000047,	PLACE2000072,
	PLACE2000097,	PLACE2000136,	PLACE2000246,	PLACE2000302,	PLACE2000379,	PLACE2000394,
5	PLACE2000425,	PLACE2000427,	PLACE2000477,	PLACE3000009,	PLACE3000070,	PLACE3000142,
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	PLACE3000322,	PLACE3000331,	PLACE3000352,	PLACE3000401,	PLACE3000413,	PLACE3000425,
	PLACE3000477,	PLACE4000009,	PLACE4000049,	PLACE4000089,	PLACE4000100,	PLACE4000247,
	PLACE4000250,	PLACE4000252,	PLACE4000300,	PLACE4000344,	PLACE4000367,	PLACE4000465,
10	PLACE4000489,	PLACE4000638,	SKNMC1000013,	THYRO1000017,	THYRO1000026,	THYRO1000034,
	THYRO1000072,	THYRO1000132,	THYRO1000173,	THYRO1000190,	THYRO1000197,	THYRO1000221,
	THYRO1000253,	THYRO1000270,	THYRO1000279,	THYRO1000327,	THYRO1000394,	THYRO1000438,
	THYRO1000558,	THYRO1000569,	THYRO1000585,	THYRO1000596,	THYRO1000625,	THYRO1000637,
	THYRO1000676,	THYRO1000734,	THYRO1000777,	THYRO1000783,	THYRO1000805,	THYRO1000843,
15	THYRO1000934,	THYRO1001033,	THYRO1001347,	THYRO1001405,	THYRO1001411,	THYRO1001534,
	THYRO1001573,	THYRO1001584,	THYRO1001602,	THYRO1001605,	THYRO1001772,	THYRO1001854,
	VESEN1000122,	Y79AA1000037,	Y79AA1000065,	Y79AA1000181,	Y79AA1000231,	Y79AA1000349,
	Y79AA1000355,	Y79AA1000368,	Y79AA1000538,	Y79AA1000782,	Y79AA1001023,	Y79AA1001145,
	Y79AA1001391,	Y79AA1001541,	Y79AA1001585,	Y79AA1001705,	Y79AA1001781,	Y79AA1001923,
20	Y79AA1001963, Y	79AA1002125, Y79	AA1002229, Y79AA	1002407, Y79AA10	02487.	
	[0244] These clo	nes are ultraviolet r	adiation damage-as	sociated clones.		

Table 12

Expression of each cDNA in human tissues (containing clones that are not described in Examples.)

				9 L	Barria V	Widon T	11	Spleen
lone_name	Heart	Lung	P. gland	Thymus	Brain	Kidney	Liver	
APDH (Cr1)	38. 210	32.670		13.580	11.230	21.120	24. 910	22.440
8 actin(Cr2)	279.280	368.870	111.100 1	17.500	92.880	114.650	82.990	256.790
DRGL1000005	53.882	23,005	32.749	22.858	26. 564	24. 940	22.644	27.001
WRGL1000007	94.778	85. 185	160, 457	67, 191	101.768	62, 489	67. 150	73. 543
DRGL1000009	11. 141	50. 520.	10. 357	7.177	6.013	5. 219	14. 272	21. 225
			29. 358	19.473	24.898	30.747	49. 220	22. 221
ADRGL1000011	71.656	24.579					19.658	11.288
ADRGL1000027	36.238	25. 252	20.855	7. 328	11.195	14. 298		
ADRGL1000058	66. 209	129.497	55. 226	49.241	30.219	55.872	67.027	243.436
ADRGL1000069	38.630	23.459	28. 991	12.540	27.353	33.633	28.774	20.911
ADRGL1000077	97.465	63.656	448. 427	83.412	71, 108	53.740	67. 906	89. 439
ADRGL1000092	89. 423	45. 692	55.810	26.033	44, 148	73.339	96.037	73.09
ADRGL 1000099	73.675	24. 424	36. 128	17.024	25. 964	41.391	42.837	29,666
ADRGL1000135	141.745	63.974	77.017	24.777	33.549	58. 986	295, 009	84. 98
ADRGL 1000147	394. 563	155. 829	271.210	92.899	165. 627	251.266	253. 420	150. 294
			39. 296	15. 194	16. 125	20.040	33.720	23. 271
ADRGL1000159	50.073	25. 425				27.027	47, 169	20.71
ADRGL1000160	69. 386	31.051	59. 416	20.154	39. 799			
ADRGL1000171	57.047	23.011	43.063	23.860	40. 581	59.814	117. 055	32.63
ADRGL1000181	45. 892	18.666	34. 476	15.434	34. 225	32.962	39.693	16. 33
BGG111000015	153. 242	42.337	92.865	41.003	45. 168	88. 524	85. 990	73.39
BGG111000016	177.367	94.731	119.688	34.159	30.249	98.806	98.783	39. 20
BGG111000017	84.712	32.614	38, 131	20.878	18.769	32.340	39.666	20.75
BGG! 11000022	52. 468	20. 452	67.167	12.167	11.158	18.241	19, 197	11.93
		17.072	40. 883	12.585	13.313	15, 525	16. 757	13, 40
BGG111000031	30.008		51.176	26.964	43. 122	43.770	49, 107	38.77
8GG111000042	49. 926	36. 336		20.304	12.650	25. 784	18, 430	25, 38
BGG111000046	31.618	26.472	34, 182	31.854				
	5031.103	2993. 496			6973.542			3649.14
BNGH41000025	91.717	35.026	73. 901	27.713	30.765	36.523	37.596	47.07
BNGH41000026	176,757	77. 439	98. 345	35.807	56.991	91.310	75.797	70.24
BNGH41000027	65.029	56. 353	25.896	22.494	12.763	23.748	17.836	23.85
BNCH41000035	148.779	66.776	119.727	56.576	60.996	96.959	72.461	64.45
BNGH41000037	79.500	29.611	43. 438	18.317	20.857	36.272	27. 525	24.77
BNGH41000042	224. 484	110.084		104. 351	102.259	125. 323	86.783	122.95
BNGH41000048	56.144	32. 253	54.063	14.729	27.312	22.435	29. 566	28. 93
BNGH41000056	67. 258	18.694	30.075	15.602	10.072	20. 735	16, 100	7.64
		46, 173	77.657	35. 329	40.900	50.029	50.841	45. 28
BNGH41000087	98. 262				5. 469	22.794	10.725	12.41
BNGH41000091	50.895	16.985	28. 392	10.147	27.072		25. 410	24. 9
BNGH41000157	69.043	34. 730	40. 597	18.088		22.074		25.84
BNGH41000169	44.850	21.770	28.655	11.403	25.991	28.509	25.634	
BNGH41000181	17. 163	15, 689	13.948	3.996	9, 287	13.139	15, 553	16. 57
BNGH41000198	81.510	36.250	60.860	20.585	25.929	35.751	31.695	28. 37
BNCH41000219	30. 302	25. 156	22. 187	13.757	11.208	15. 235	27. 285	35.70
BNGH41000229	252.790	65, 948	93.499	51, 108	92.555	101.245	96.716	78. 2
BNGH41000237	85.757	46, 997	55. 170	26.780	33.764	47.456	37.007	39.1
BNGH41000238	17.744	36.938	42.360	14, 922	35.749	42.848	39. 238	13.2
BNGH41000243	45.446	23.667	44, 798	20.875	10.516	23.918	22. 443	27.0
BNGH41000270	60.889	18, 651	29.618	10.724	15.979	12.351	19. 152	22.3
BRAWH1000004	43.673	28. 539	7.640	11.388	19, 198	14. 903	32. 353	23.7
BRAWH1000018	59, 409	17. 941	102. 270	17, 107	709.078	25.732	24. 214	24.7
				21.042			66.754	27. 9
BRAWH1000021	104.772	29. 951	51.142		1169.154	70.731	79.670	40. 9
BRAWH1000027	152. 205	47.310	67.089	32.199	64. 521		64. 886	
BRAWH1000029	106.376	49. 221	55. 840	40.856	59, 552	56.487		100.1
BRAWH1000040	29.419	16.761	31, 101	16.622	30,633	18.200	17. 998	15.1
BRAWH1000050	161.264	71.786	118.976	51.863	61.542	97.720	81.271	69. 1
BRAWH1000051	74.067	34. 341	44.047	20.726	30.434	42.055	53. 856	24.6
BRAWH1000060	68, 789	22.598	35.012	16.493	19. 127	38.662	34. 923	28.0
BRAWH1000075	17, 318	16.898		8.901	18.133	17.219	9. 321	11.2
BRAWH1000081	43.025			7.655	123.677	17.673		9.8
BRAWH1000084	174. 384				152. 188		110.167	102.2
					116. 975		746. 700	35. 9
BRAWH1000095	118. 239				145.013			
	146.112							63.
BRAWH1000096					1 PM /11()	1 /3./07	63.827	
BRAWH1000097	95. 841	72. 506					10 000	77.7
	95. 841 11. 943 134. 838	19.037	18. 950	13.536	92.145 96.396	16.582		10. 2 57. 3

Table 13

BRAWH1000104	25, 414 T	18. 303	14. 825	7.695	38.918	23. 970	23.794	11.048
BRAWH1000107			12.463	5.518	6. 355	5. 084	9. 107	6. 573
BRAWH1000110	16.949	5.616 492.704		383.612	368, 156	369, 621	277. 348	340, 450
	615. 476		92.209				90. 369	57. 301
BRAWH1000111	175. 556	68. 459		45.974	64.703 57.720	81.723	87.016	35.866
BRAWH1000135	199. 303	38.098	72.093	26.809		91.668		
BRAWH1000190	56, 386	41.640	57.914	22.782	55.671	40. 034	35. 280	40.134
HEMBA 1000005	11.985	23. 427	18.882	9.766	12.656	9. 959	23. 443	21.677
HEMBA 1000006	37.398	24. 521	24. 529	15. 587	22.317	13.336	16.038	15. 295
HEMBA1000012	81.820	57.193	66.828	26.683	55. 423	58. 731	85.614	66. 259
HEMBA 1000020	157.967	64.157	115.635	51.940	77.293	77. 321	83.989	74. 362
HEMBA1000030	82.882	35. 447	66.058	26.464	40.990	60.871	47.058	50.652
HEMBA1000034	47.434	17.878	50.696	5.594	14.005	6.673	24.652	7.134
HEMBA1000042	147.376	94.003	330.908	69.071	76.472	55. 477	37.783	60.479
HEMBA 1000045	28.478	20. 289	20.548	12.445	11.835	22.788	11.196	15.775
HEMBA1000046_	85.160	84.475	242.940	57.017	68.488	45. 288	37.098	47.486
HEMBA1000047	21.380	18.899	18.166	11.393	11.185	12. 292	6.491	12.018
HEMBA 1000048	243.559	55.114	84. 448	24. 247	43.131	99, 333	57.041	37. 362
HEMBA 1000050	22.711	11.876	21.972	7.477	4.096	13.675	10.347	7.770
HEMBA1000053	45.071	26.410	38.158	15.982	30.754	36.740	34. 184	24. 269
HEMBA1000060	101.197	34. 766	50.643	19.938	34.641	54.061	42.309	22.530
HEMBA1000072	240.166	213.938	224.688	163.030	115.246	207.809	112.361	276.098
HEMBA1000073	23. 202	9. 580	10.815	1.698	6. 680	18. 155	12.304	14. 973
HEMBA 1000076	95. 997	46.783	177.931	32.617	48.964	50. 792	33.947	44. 142
HEMBA1000084	66.603	25.710	48. 434	18.006	22.553	38.118	40.479	29.683
HEMBA1000087	70.084	17.515	26.544	8. 450	17. 590	29. 220	19.519	22. 565
HEMBA1000088	15.474	8.614	19.903	4.775	4.519	11.446	34. 905	6. 528
HEMBA1000091	80.622	38.604	59.393	23.956	44, 939	49.760	33. 946	24.614
HEMBA1000111	85.814	95.270	270.642	75. 147	54. 384	70.071	29. 529	55. 422
HEMBA1000121	55.476	43.368	146.465	37.419	29. 398	30.694	17.702	30.398 23.965
HEMBA1000128	37.278	27.165	34.516	13.619	17.702 27.243	28.069	12.834	26.851
HEMBA 1000129	51.488	19.659	44. 907 32. 107	14.353	13.502	30. 959 11. 152	8. 907	20.635
HEMBA1000141 HEMBA1000146	12.961	24.515 11.479	20.418	8. 202	9.575	14.877	10,000	7.817
	29.273 534.562	326. 814	684. 147	211.774	218.448	322. 246	235.752	256.883
HEMBA1000150 HEMBA1000154	95. 272	92.253	101.483	54.276	42.896	75. 526	92.689	188.019
HEMBA1000156	50.177	72. 591	58.026	31.149	21.865	38. 964	27.634	50. 220
HEMBA 1000158	260.718	63. 920	89.680	36.337	44.915	93. 421	111.344	53. 562
HEMBA1000168	74.415	61.152	62.826	30.512	23. 287	34. 966	44.005	33. 564
HEMBA1000180	28.502	22. 412	28.571	11.701	19. 230	10.903	11.731	14, 102
HEMBA1000185	115.723	50.661	213.994	51.166	43. 435	56. 261	38.862	44. 992
HEMBA1000188	21. 302	14. 879	16. 948	11. 392	11.821	10.656	12.501	6. 979
HEMBA1000193	14. 122	8.318	11.905	7.519	4.736	3. 349	8, 544	7.842
HEMBA1000194	54.688	49. 534	143.817	37.736	20. 221	34. 328	23.359	56. 497
HEMBA1000201	21.062	14.098	8.690	6.237	5, 109	5.059	9.317	10.522
HEMBA1000213	22.388	25. 532	25,777	8. 470	17.320	9.084	8.469	11.766
HEMBA1000216	65.935	51.368	92.680	19. 202	33.659	40. 971	36.328	34.891
HEMBA1000227	52.577	31.332	34.925	19.503	18.411	21.504	22.590	25. 781
HEMBA1000231	114.369	54. 299	131.256	38.550	43.246	29.778	24. 266	30, 410
			1 100 220	58. 292	93.250	57,000	49.319	59, 288
HEMBA1000237	91.024	91, 360	199.338	30. 232	33.230			
HEMBA1000243	53.456	43.969	117.519	38.431	25, 396	32,604	38, 910	32.153
HEMBA1000243	53.456	43.969			25, 396	32,604		32.153
		43.969	117.519	38.431		32,604	38, 910	32.153
HEMBA1000243 HEMBA1000244	53.456 173.469	43.969 104.733	117.519 115.584 14.367 130.009	38. 431 33. 079	25, 396 65, 527 16, 095 37, 481	32.604 124.532 13.221 24.002	38. 910 90. 927 11. 516 20. 553	32.153 78.610 11.018 13.215
HEMBA 1000243 HEMBA 1000244 HEMBA 1000251	53.456 173.469 22.709	43.969 104.733 12.333	117.519 115.584 14.367	38. 431 33. 079 9. 019 20. 848 9. 473	25, 396 65, 527 16, 095	32.604 124.532 13.221	38. 910 90. 927 11. 516 20. 553 13. 690	32. 153 78. 610 11. 018
HEMBA1000243 HEMBA1000244 HEMBA1000251 HEMBA1000254 HEMBA1000264 HEMBA1000269	53. 456 173. 469 22. 709 74. 060 29. 478 36. 718	43. 969 104. 733 12. 333 35. 626 15. 248 13. 465	117.519 115.584 14.367 130.009	38. 431 33. 079 9. 019 20. 848	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705	32.604 124.532 13.221 24.002 11.228 12.833	38. 910 90. 927 11. 516 20. 553 13. 690 7. 348	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793
HEMBA 1000243 HEMBA 1000244 HEMBA 1000251 HEMBA 1000254 HEMBA 1000264 HEMBA 1000269 HEMBA 1000275	53.456 173.469 22.709 74.060 29.478 36.718 66.201	43.969 104.733 12.333 35.626 15.248 13.465 39.367	117.519 115.584 14.367 130.009 23.537 28.932 84.077	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871	32.604 124.532 13.221 24.002 11.228 12.833 49.267	38, 910 90, 927 11, 516 20, 553 13, 690 7, 348 36, 211	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871
HEMBA1000243 HEMBA1000244 HEMBA1000251 HEMBA1000254 HEMBA1000269 HEMBA1000275 HEMBA1000275	53.456 173.469 22.709 74.060 29.478 36.718 66.201 33.299	43.969 104.733 12.333 35.626 15.248 13.465 39.367 36.073	117.519 115.584 14.367 130.009 23.537 28.932 84.077 54.357	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751	38, 910 90, 927 11, 516 20, 553 13, 690 7, 348 36, 211 21, 696	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871 30. 785
HEMBA 1000243 HEMBA 1000244 HEMBA 1000251 HEMBA 1000254 HEMBA 1000264 HEMBA 1000269 HEMBA 1000275	53.456 173.469 22.709 74.060 29.478 36.718 66.201	43.969 104.733 12.333 35.626 15.248 13.465 39.367	117.519 115.584 14.367 130.009 23.537 28.932 84.077	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384	38. 910 90. 927 11. 516 20. 553 13. 690 7. 348 36. 211 21. 696 56. 916	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871 30. 785 92. 414
HEMBA1000243 HEMBA1000244 HEMBA1000251 HEMBA1000254 HEMBA1000269 HEMBA1000275 HEMBA1000275	53.456 173.469 22.709 74.060 29.478 36.718 66.201 33.299 93.815 12.439	43.969 104.733 12.333 35.626 15.248 13.465 39.367 36.073 121.083 24.935	117.519 115.584 14.367 130.009 23.537 28.932 84.077 54.357 171.037 29.793	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017 123. 971 37. 925	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384 9.632	38, 910 90, 927 11, 516 20, 553 13, 690 7, 348 36, 211 21, 696 56, 916 2, 866	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871 30. 785 92. 414 7. 311
HEMBA 1000243 HEMBA 1000244 HEMBA 1000251 HEMBA 1000254 HEMBA 1000264 HEMBA 1000269 HEMBA 1000275 HEMBA 1000280 HEMBA 1000280	53.456 173.469 22.709 74.060 29.478 36.718 66.201 33.299 93.815	43. 969 104. 733 12. 333 35. 626 15. 248 13. 465 39. 367 36. 073 121. 083 24. 935 30. 009	117.519 115.584 14.367 130.009 23.537 28.932 84.077 54.357 171.037 29.793 145.363	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720 93. 484 10. 840 25. 471	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017 123. 971 37. 925 9. 769	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384 9.632 16.272	38. 910 90. 927 11. 516 20. 553 13. 690 7. 348 36. 211 21. 696 56. 916 2. 866 9. 701	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871 30. 785 92. 414 7. 311 15. 510
HEMBA1000243 HEMBA1000244 HEMBA1000251 HEMBA1000254 HEMBA1000264 HEMBA1000275 HEMBA1000275 HEMBA1000282 HEMBA1000282	53. 456 173. 469 22. 709 74. 060 29. 478 36. 718 66. 201 33. 299 93. 815 12. 439 45. 269 14. 803	43. 969 104. 733 12. 333 35. 626 15. 248 13. 465 39. 367 36. 073 121. 083 24. 935 30. 009 5. 750	117. 519 115. 584 14. 367 130. 009 23. 537 28. 932 84. 077 54. 357 171. 037 29. 793 145. 363 10. 615	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720 93. 484 10. 840 25. 471 5. 725	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017 123. 971 37. 925 9. 769 2. 559	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384 9.632 16.272 8.602	38. 910 90. 927 11. 516 20. 553 13. 690 7. 348 36. 211 21. 696 56. 916 2. 866 9. 701 8. 358	32, 153 78, 610 11, 018 13, 215 3, 797 24, 793 38, 871 30, 785 92, 414 7, 311 15, 510 9, 224
HEMBA1000243 HEMBA1000244 HEMBA1000251 HEMBA1000254 HEMBA1000269 HEMBA1000275 HEMBA1000280 HEMBA1000280 HEMBA1000288 HEMBA1000287 HEMBA1000288 HEMBA1000288 HEMBA1000288	53. 456 173. 469 22. 709 74. 060 29. 478 36. 718 66. 201 33. 299 93. 815 12. 439 45. 269 14. 803 27. 085	43. 969 104. 733 12. 333 35. 626 15. 248 13. 465 39. 367 36. 073 121. 083 24. 935 30. 009	117. 519 115. 584 14. 367 130. 009 23. 537 28. 932 84. 077 54. 357 171. 037 29. 793 145. 363 10. 615 21. 195	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720 93. 484 10. 840 25. 471 5. 725 9. 790	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017 123. 971 37. 925 9. 769 2. 559 16. 909	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384 9.632 16.272 8.602 12.402	38. 910 90. 927 11. 516 20. 553 13. 690 7. 348 36. 211 21. 696 56. 916 2. 866 9. 701 8. 358 15. 289	32, 153 78, 610 11, 018 13, 215 3, 797 24, 793 38, 871 30, 785 92, 414 7, 311 15, 510 9, 224 17, 159
HEMBA 1000243 HEMBA 1000244 HEMBA 1000251 HEMBA 1000254 HEMBA 1000264 HEMBA 1000269 HEMBA 1000280 HEMBA 1000282 HEMBA 1000282 HEMBA 1000288 HEMBA 1000288 HEMBA 1000288 HEMBA 1000298 HEMBA 1000298 HEMBA 1000298 HEMBA 1000296	53. 456 173. 469 22. 709 74. 060 29. 478 36. 718 66. 201 33. 299 93. 815 12. 439 45. 269 14. 803 27. 085 98. 491	43. 969 104. 733 12. 333 35. 626 15. 248 13. 465 39. 367 36. 073 121. 083 24. 935 30. 009 5. 750 22. 625 119. 119	117. 519 115. 584 14. 367 130. 009 23. 537 28. 932 84. 077 54. 357 171. 037 29. 793 145. 363 10. 615 21. 195 304. 884	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720 93. 484 10. 840 25. 471 5. 725 9. 790 73. 660	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017 123. 971 37. 925 9. 769 2. 559 16. 909 85. 595	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384 9.632 16.272 8.602 12.402 48.175	38, 910 90, 927 11, 516 20, 553 13, 690 7, 348 36, 211 21, 696 56, 916 2, 866 9, 701 8, 358 15, 289 43, 496	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871 30. 785 92. 414 7. 311 15. 510 9. 224 17. 159 66. 547
HEMBA1000243 HEMBA1000244 HEMBA1000251 HEMBA1000254 HEMBA1000269 HEMBA1000275 HEMBA1000280 HEMBA1000280 HEMBA1000288 HEMBA1000287 HEMBA1000288 HEMBA1000288 HEMBA1000288	53. 456 173. 469 22. 709 74. 060 29. 478 36. 718 66. 201 33. 299 93. 815 12. 439 45. 269 14. 803 27. 085	43. 969 104. 733 12. 333 35. 626 15. 248 13. 465 39. 367 36. 073 121. 083 24. 935 30. 009 5. 750 22. 625	117. 519 115. 584 14. 367 130. 009 23. 537 28. 932 84. 077 54. 357 171. 037 29. 793 145. 363 10. 615 21. 195	38. 431 33. 079 9. 019 20. 848 9. 473 20. 412 38. 846 24. 720 93. 484 10. 840 25. 471 5. 725 9. 790	25. 396 65. 527 16. 095 37. 481 3. 863 9. 705 77. 871 38. 017 123. 971 37. 925 9. 769 2. 559 16. 909	32.604 124.532 13.221 24.002 11.228 12.833 49.267 35.751 70.384 9.632 16.272 8.602 12.402	38. 910 90. 927 11. 516 20. 553 13. 690 7. 348 36. 211 21. 696 56. 916 2. 866 9. 701 8. 358 15. 289	32. 153 78. 610 11. 018 13. 215 3. 797 24. 793 38. 871 30. 785 92. 414 7. 311 15. 510 9. 224 17. 159 66. 547

Table 14

					44.4	20 100	A. A	44 010	40 071
	HEMBA1000303	129. 286	51.013	88.777	32.513	50.462	82. 994	44.818	49. 271
	HEMBA 1000304	112.022	67.470	328.677	54.678	79. 305	43. 526	38. 469	55. 762
	HEMBA1000307	14.054	22.013	31.964	13, 167	15, 571	7.974	10.014	8.685
5			69.330	183. 923	45. 322	45.087	52, 968	37.741	38. 246
	HEMBA 1000312	97.082							
	HEMBA 1000318	16. 164	16.264	18.766	11.688	3.620	10.732	8. 295	14.675
	HEMBA1000327	29. 404	59.618	81.347	41.731	85.004	48. 526	49. 421	46.866
	HEMBA1000333	16. 964	13.930	14.530	1.872	5.776	1. 571	0.392	3.743
	HEMBA1000338	121.878	62.572	348.751	55.463	49.114	38, 561	30.698	40.644
						5. 872	16. 551	10, 139	14. 088
10	HEMBA 1000343	25. 229	29. 781	46. 395	20.673				
	HEMBA1000349_	23.061	12.586	31.755	7.020	17.658	11. 522-	14.807	15.611
	HEMBA 1000351	92.847	57. 338	196.577	41.762	37.094	35. 370	27. 545	28. 615
	HEMBA 1000355	85.210	38, 388	64. 299	18.101	33, 114	43.511	37.808	26.628
		60.438	38.786	62, 442	20.784	17.694	38.058	40. 431	28. 899
	HEMBA 1000356					42. 793	39. 432	26.076	44. 579
	HEMBA1000357	84.898	55. 990	206.803	54. 151				
45	HEMBA1000366	47. 131	42.031	90. 450	27.056	20.718	23.499	14.632	23. 547
15	HEMBA1000369	71.428	40.685	54. 384	17.613	21.422	34. 985	37.622	36.900
	HEMBA1000370	16.354	14.949	22.988	7.916	18.390	15. 359	13.426	5.647
	HEMBA1000376	80, 183	75. 300	201.705	55. 266	66.687	44.612	55. 386	56.070
							74.007	57.619	79.876
	HEMBA1000387	100.497	129. 367	351.196	80. 257	104. 250			
	HEMBA 1000389	69. 342	34.021	71.118	22.346	27. 319	47. 936	53.026	34. 161
	HEMBA 1000390	19. 206	25.788	21.028	12.401	18. 372	13.751	16.243	15.036
20	HEMBA1000392	19.400	22.884	44, 179	8.776	11.742	10.594	12.266	12.463
	HEMBA1000396	75. 409	50. 195	81.870	27.979	30.393	31, 235	17.771	19. 584
				25. 987	10.341	31.398	31. 214	50, 056	18. 580
	HEMBA1000411	35. 966	24. 397						
	HEMBA 1000418	8. 165	10.778	14.987	4. 031	12.495	7. 913	6. 363	2. 306
	HEMBA 1000422	93.699	38. 329	85. 266	39. 826	45. 992	44. 729	42.886	34, 308
	HEMBA 1000428	51.017	30.690	79.229	26.579	24. 840	17. 767	18.424	18.608
25	HEMBA 1000434	1.747	3.214	11, 346	1.210	1.602	2. 927	2.788	2.756
			7.698	16. 227	7. 252	3. 336	17. 969	11.723	10. 545
	HEMBA1000442	21.750					31.691	47.856	102. 287
	HEMBA 1000443	67. 291	35.910	34.775	26.420	16.860			
	HEMBA 1000446	236.986	69.546	90. 283	32.233	34.107	119. 377	108, 645	60. 266
	HEMBA1000456	95.368	37.560	63.451	22.640	41.092	65, 256	62.972	43. 493
	HEMBA 1 000459	28.924	35. 333	74. 945	20.475	25.324	26. 253	13.654	31.317
30	HEMBA1000460	18. 649	27.246	21.973	9.513	15.230	14, 091	9.746	16.955
30					31.332	83.626	109. 503	92.971	62. 126
	HEMBA 1000462	220. 184	42.636	96.490					
	HEMBA 1000464	34. 277	15. 137	27.210	10.862	15. 595	20, 793	16.716	16. 539
	HEMBA1000468	41.755	41.852	68.356	10,400	23.452	43, 909	24.048	22.968
	HEMBA1000469	68. 229	71.011	256.705	47.636	29.853	34. 188	22.568	39.190
	HEMBA1000477	185. 220	47.546	102.939	26.276	40.188	95, 247	52.454	28. 109
			37. 528	24.407	17, 115	24. 182	29.826	20.717	25.819
35	HEMBA1000481	47.276							
	HEMBA 1000488	96. 226	31.249	71.522	21.667	27.715	44, 499	53.708	33.306
	HEMBA1000490	29.915	13.747	32.568	14.002	12.056	6. 900	11.274	7.559
	HEMBA1000491	80.198	22.903	47.786	20.675	32.551	52.682	37.109	28. 282
	HEMBA1000498	191.186	112,767	454. 998	88.614	102.997	82.927	53.205	120.837
	HEMBA 1000501	57. 318	55. 923	180.158	44. 170	27.291	34, 954	18.532	34.117
			5. 893	7, 152	1.726	0.520	2. 245	2.551	1.091
40	HEMBA 1000504	1.033						19.508	31.925
	HEMBA1000505	55. 746	36.631	48, 155	21.562	14.691	34. 729		
	HEMBA 1000507	204. 165	114.530	305. 249	86.138	81.505	97. 289	230.331	95. 150
	HEMBA 1000508	205.724	105.067	309.791	72.709	70.180	77. 388	63.849	45. 940
	HEMBA1000518	39. 157	29, 100	31,505	16.650	14.796	15.847	24.729	17.601
	HEMBA1000519	166. 937			148.478	123.978	128. 646	85.670	111.078
						10.619	1. 488	9, 513	9. 395
45	HEMBA 1000520	0.000	0.000	0.000					
	HEMBA 1000523	38, 708	22.090	40.875	13.852	21.603	32. 384	20.478	21. 422
	HEMBA1000531	21.874	34.044	40.027	12.264	11.034	29.775	20.421	12.540
	HEMBA1000534	0.000	0.000	0.000	34.434	48.940	25. 365	41.242	72. 583
	HEMBA1000538	0.000	0.000	0.000		19.981	17.606	26.698	23. 904
				33.145	42.629	27.059	33. 931	16.639	31.893
	HEMBA1000540	21.974	47.343						
50	HEMBA 1000542	64.656	33. 152	58.093		35. 278	55. 508	47.917	47.623
50	HEMBA1000545	148.870	136.401	48.802	8. 499	12.534	7, 119	25.484	15.094
	HEMBA 1000547	14.825	20, 199	32.694		22.359	12.020	13.535	20. 227
	HEMBA1000551	163.806	171.089	543.876		115.775	116.646	69.596	152.516
	HEMBA 1000555		20, 199	25. 801	24. 488	14.071	15. 431	5. 986	10. 933
		10.531					31.872		30. 926
	HEMBA1000557	80.051	48. 396	168.724					
	HEMBA 1000561	56. 992	22.797	51.047	10. 187	16.301	34.904	24.661	22.470
55									

Table 15 ·

	TUTUDA LODGE ES		14 5 76						
	HEMBA1000563	9.473	11.545	18.205	6.139	12.689	10.132	7.939	14. 253
	HEMBA1000567	41.385	38.483	27.881	32. 207	15. 544	26.052	15.086	86.601
	HEMBA1000568	44.686	33.379	126.524	26. 300	22, 533		26.970	
5							17.402		18.707
	HEMBA1000569	58, 184	27.187	41,012	21.787	12, 925	36. 191	33.944	23. 225
	HEMBA1000575	155.833	155.759	434. 526	92.140	79, 143	69.949	59. 928	71.189
	HEMBA1000588	41.087	26.072	31.610	14. 580				
						18,024	18. 458	23. 553	13. 279
	HEMBA1000590	29.693	17.090	23.618	7.069	6.633	16.725	20.068	13.042
	HEMBA1000591	106.772	54.874	98.079	34.099	31,776	57, 170	48, 488	32.766
	HEMBA1000592	7.408	10.031	9.435	9. 551	8, 209	5. 142	7.480	
10	HEMBA1000594								10.319
		18, 401	11.048	22.547	15. 327	9. 596	12.099	8.751	6.852
	HEMBA1000604	96.047	78.462	146.030	49.571	36.099	70.815	41.797	47.748
	HEMBA1000607	46.819	15.606	46.037	9. 438	19, 149	21.038	17.317	25. 404
	HEMBA1000608	8, 985	3.040	6.705	0.000	7. 378	4. 453	0.000	5. 544
	HEMBA1000622	45. 570	55.746	113.666	40.310	18. 167	19.390	15.895	29. 149
15	HEMBA1000634	126, 532	49.146	138.073	29.094	95.787	79.662	60.271	71.657
	HEMBA1000636	151.899	51.270	126.200	39, 161	51.864	62.611	54.056	39. 415
	HEMBA1000637	33, 241	23.587	39. 380	18.047	16.265	30.075		
								28. 226	24. 559
	HEMBA1000655	80.165	70.766	219. 283	58. 901	61.320	45. 821	40.741	62. 639
	HEMBA1000657	60,961	31.993	41.401	18.008	30.565	35. 201	35.611	42. 178
	HEMBA1000652	8,600	8.490	11.263	5. 475	2.201	6.140	1.557	2. 504
	HEMBA1000664	14. 358	5. 082	3.637	2.670	3.516			
20							4.913	3.094	3. 579
	HEMBA1000671	11.588	15.473	26.067	17.940	8.865	7.647	10.779	21.196
	HEMBA1000673	73.174	77.410	193. 253	46.051	34. 388	33.975	25.896	31.646
	HEMBA1000675	7.666	12.047	22. 123	5.764	42.036	15.788	10.254	15. 555
	HEMBA1000678	7.453	12.314	21.083	12.174	14.897	12.628		
	HEMBA1000682							6.969	6. 584
		118, 965	125.696	255. 731	86.894	61.443	66.299	49.060	82. 939
25	HEMBA1000686	25.079	17.463	23. 126	12.722	10.282	13.835	21.393	18. 154
20	HEMBA1000702	206.683	94. 357	266.585	62.386	79.930	90, 914	98.397	60. 559
	HEMBA1000705	25. 430	25. 862	47. 190	13.191	19.599	26.364	25.013	
	HEMBA1000713								18.833
	DEMONITURE 13	56. 893	25. 288	70.751	17.660	24. 138	23.311	21.805	21.736
	HEMBA1000718	50.149	43.869	128. 515	28. 289	23.213	18. 458	10.003	17.419
	HEMBA1000719	37.969	17.467	28. 513	12.147	12.768	22. 543	14.744	14. 432
	HEMBA1000722	15.150	9.762	14.699	6.768	11.726	12.080	5. 907	9. 953
30	HEMBA1000726	159.817	111.276	463. 937					
					91.448	109.093	58. 587	46.517	70.087
	HEMBA1000727	22.867	26.803	28. 886	21.475	11.199	14.966	8.634	30.401
	HEMBA1000732	28. 530	11.011	12.790	4.617	3. 548	13.325	19.978	13.472
	HEMBA1000736	24. 568	21.982	21.410	7.431	11.376	41.026	31.698	16.801
	HEMBA1000743	0.741	4. 467	1.793	1.637	1.227			
							3.642	4.563	3.368
35	HEMBA1000745	8. 930	7.067	14. 546	3.314	10.067	5. 403	9. 225	6.085
00	HEMBA1000747	21.442	12.487	25.662	17.081	5. 384	10. 287	9.865	8.267
	HEMBA1000748	22. 924	14.885	35. 721	12.634	3.045	11.508	4.110	11.756
	HEMBA1000749	67. 267	50.825	159.211	43.879	20. 345	29.613	19.447	
	HEMBA1000752	54, 929	35.778	162.005					31.693
					28. 209	31.540	25, 132	15.650	20.776
	HEMBA1000753	120.889	83.878	155.892	48.092	54. 307	53. 238	38.941	39. 331
	HEMBA1000757	20. 234	22. 592	52.608	29. 935	23.071	24. 503	14. 548	43.779
40	HEMBA1000760	12.599	38.665	19.973	15.800	30.188	14. 155	10.570	39. 229
	HEMBA1000769	114.956	74. 924	304. 424	66.815	39. 365	48. 405	39.918	55. 931
	HEMBA1000773	2. 162	5. 360						
				11.883	4, 445	0.965	3.158	3. 956	2.663
	HEMBA1000774	128. 563	115.732	330, 111	84.461	69.618	59.363	42.656	56. 152
	HEMBA1000780	6.850	7.130	24.176	6.924	6.903	6.546	6.667	9.576
	HEMBA1000783	8. 127	5.076	13.701	3. 276	8.863	5.241	5. 435	4. 429
45	HEMBA1000791	41.433	51.546	108.542	29.633				
						42.735	44. 515	43. 187	40.856
	HEMBA1000793	108, 761	30, 885	54. 568	18.670	31.512	54.669	45. 458	34. 788
	HEMBA1000802	15.062	11. 125	9.052	10.300	11.505	12. 950	15. 354	16. 952
	HEMBA1000813	106.763	52.683	69.701	32.507	44. 369	65.862	59.842	56.799
	HEMBA1000817	19. 480	7.070	17.915	4, 016				
						15. 239	18.434	11.273	8.079
E0	HEMBA1000822	9. 520	10.358	15.760	7.218	8.704	11.185	6.639	4.662
50	HEMBA1000827	96.001	12.420	24.041	8, 305	24.000	6.709	3. 488	8. 591
	HEMBA1000833	53, 675	28.970	35.897	14, 604	26. 383	29.036	20. 591	14, 341
	HEMBA1000835	74.696	67.353	83.737	34, 349	42. 834	61.145		
		74. 227						66.784	52.015
	HEMBA1000843		54. 197	92.042	37, 825	58. 573	98. 943	87.569	55. 077
	HEMBA1000851	23.913	14.070	13.081	6.847	8.634	12.419	19, 200	22. 286
	HEMBA1000852	56.702	54.074	105.085	31, 127	34. 200	31.843	28.843	30. 311
55	HEMBA1000867	15. 548	10.247	11.912	6.256	1. 227	12.374		
		1 10. 070	1 .0.671	1 11.312	. 0.230	1.441	14.314	8.518	5. 611

Table 16

FIERBAIL (100965)										
HEBBA1000975		HEMBA1000869	19.696 T	18. 785	34.039	15.061	6.930	13. 298	14. 305	14.712
HEBBA1000912		HEMBA LODGETO		38 245	44 665	12 547	23 970	41 195	21 911	17 508
HEBBA1000975 35	•									
	5	HEMBA1000872	45.848	46.546	86. 933					
HENDAIDODRIF 89, 976 \$6, 654 194, 869 42, 595 \$7, 670 \$51, 567 \$6, 331 40, 884 HENDAIDODRIB 45, 409 18, 456 30, 665 12, 448 8, 114 19, 529 24, 789 16, 299 11, 592 14, 380 14, 379 17, 35, 39 3, 363 5, 327 13, 020 16, 299 11, 592 14, 380 17, 381 22, 29, 31 3, 566 67, 073 19, 383 22, 24, 789 16, 299 18, 286 7, 073 19, 383 22, 24, 789 16, 299 18, 286 7, 073 19, 383 22, 24, 789 16, 299 18, 286 7, 073 19, 383 22, 24, 789 16, 299 18, 286 7, 073 19, 383 22, 24, 789 16, 299 18, 286 7, 073 19, 383 22, 24, 789 11, 592 18, 280 1	J	HEMBA1000875	35 460	41 166	32, 238	11, 297	35, 077	29, 781	19, 453	23. 540
HEMBA1000907 22, 559 3, 656 10, 917 3, 559 3, 363 5, 327 11, 032 10, 676 HEMBA1000910 47, 107 13, 661 26, 933 5, 866 7, 073 19, 938 22, 971 11, 529 HEMBA1000918 67, 477 13, 661 26, 933 5, 866 7, 073 19, 938 22, 971 11, 529 HEMBA1000919 67, 477 13, 661 26, 933 5, 866 7, 073 19, 938 22, 971 11, 529 HEMBA1000919 67, 447 29, 880 114, 873 25, 206 16, 670 22, 595 24, 770 HEMBA1000919 67, 447 29, 880 14, 873 25, 206 16, 670 24, 539 23, 073 20, 233 HEMBA1000919 67, 467 29, 841 71, 669 12, 587 23, 834 13, 188 7, 830 13, 322 HEMBA1000915 16, 784 29, 481 71, 669 12, 587 23, 834 13, 188 7, 830 13, 322 HEMBA1000940 44, 273 87, 986 91, 786 24, 577 24, 314 31, 188 7, 830 13, 322 HEMBA1000943 14, 439 17, 702 14, 850 4, 792 8, 391 11, 855 11, 039 7, 414 HEMBA1000943 14, 439 15, 765 18, 812 29, 000 5, 772 0, 000 19, 405 9, 919 HEMBA1000950 179, 850 15, 073 44, 747 07, 719 85, 691 11, 193 82, 284 44, 494 HEMBA1000985 179, 850 15, 073 44, 747 07, 719 85, 691 11, 193 82, 284 44, 454 HEMBA1000977 51, 106 19, 281 37, 515 12, 222 19, 562 23, 744 46, 24 HEMBA1000977 50, 148 19, 281 37, 515 12, 222 19, 562 23, 744 46, 24 HEMBA1000977 50, 148 19, 281 37, 515 12, 222 19, 562 23, 874 44, 624 HEMBA1000977 50, 108 19, 281 37, 515 12, 222 19, 562 23, 874 44, 624 HEMBA1000977 50, 148 19, 281 37, 515 12, 222 19, 562 23, 874 44, 624 HEMBA1000977 50, 108 13, 532 39, 641 19, 794 27, 516 38, 200 13, 437 38, 444 HEMBA1000987 30, 47 79, 132 42, 499 15, 644 4, 099 16, 478 4, 14, 23 HEMBA1000988 12, 794 79, 132 42, 499 15, 644 4, 099 16, 478 4, 41, 52 HEMBA1000988 12, 794 79, 795 79, 795 79, 795 79, 795 79, 795 79, 795 79, 795 79, 795										
HEMBAIODOGIS		HEMBA1000907	22, 959	9. 656	10.917	3.599	3.363	5. 327	13.032	10.676
					20 665	12 448	R 174	19 529	24 789	16 299
		[HEMBA1000910]	47.107		26. 933					
		HEMBA1000918	67 437	29 880	114, 873	25, 206	16,670	25.895	26.769	24.710
HEBBAT000934	10									
HEBBAT000935										
HEBBAT000940		HEMBA1000934	162.546	35. 314	59.012	18.820		53. 492	33.824	20.798
HEBBAT000940		UENBATOOO35	16 284	29 481	71 659	12 587	23.834	13, 188	7, 830	13.322
HEMBA1000942 125.095 96.812 260.912 82.657 49.118 47.891 35.814 49.631 HEMBA1000945 13.487 12.702 14.890 4.792 8.391 11.856 1.093 7.414 HEMBA1000946 15.487 5.506 18.692 9.000 5.772 0.000 19.405 9.919 HEMBA1000950 173.860 151.073 345.747 107.319 85.595 117.093 32.928 34.494 14.8841000985 73.385 34.603 60.061 26.562 28.789 47.944 60.067 31.618 14.8841000997 50.148 19.281 7.515 12.227 19.562 28.778 7.458 7.214 4.624 14.8841000997 50.148 19.281 7.515 12.227 19.562 29.874 20.455 23.135 14.8861 19.281 19.8861 1										
THE		HEMBA I DOUG40								
HEMBAT000943		HEMBA1000942	125.095	96. 812	260.912	62.657	49.118	47. 891	35.814	49.631
HEMBAT000946 15. 461 5. 506 18. 682 9.000 5.772 0.000 19. 405 9. 919 HEMBAT000955 77. 395 34. 747 70.731 8. 559 117.093 29. 292 94. 494 HEMBAT000952 77. 395 34. 803 60.061 26. 562 28. 789 47. 944 60.067 31. 619 HEMBAT000957 50. 148 19. 281 37. 515 12. 222 19. 562 29. 374 22. 045 23. 135 HEMBAT000977 50. 148 19. 281 37. 515 12. 222 19. 562 29. 374 22. 045 23. 135 HEMBAT000977 50. 148 19. 281 37. 515 12. 222 19. 562 29. 374 22. 045 23. 135 HEMBAT000977 50. 148 19. 281 37. 515 12. 222 19. 562 29. 374 22. 045 23. 135 HEMBAT000978 51. 100 33. 450 188. 137 28. 972 24. 576 23. 735 13. 731 27. 272 HEMBAT000978 34. 417 19. 132 42. 499 15. 644 4. 009 16. 478 41. 152 14. 153 HEMBAT000981 35. 318 31. 281 38. 672 19. 544 4. 009 16. 478 41. 152 14. 153 HEMBAT000981 35. 318 31. 281 38. 672 19. 544 4. 009 6. 478 41. 152 41. 253 HEMBAT000985 7. 290 20. 183 32. 497 4. 058 6. 443 9. 035 7. 852 3. 257 HEMBAT000986 72. 707 55. 780 56. 633 22. 497 4. 058 6. 443 9. 035 7. 852 3. 257 HEMBAT000991 72. 707 55. 780 56. 717 34. 676 32. 494 41. 317 23. 483 37. 846 HEMBAT001007 72. 650 42. 553 58. 807 23. 255 34. 284 41. 317 23. 483 37. 846 HEMBAT001007 72. 650 42. 553 58. 807 23. 255 34. 284 47. 77. 17. 22. 655 HEMBAT001009 73. 843 29. 269 35. 135 10. 555 77. 15 77. 79. 17. 17. 17. 17. 17. 17. 17. 17. 17. 17				12 702		A 792	8 391	11 856	11 039	7 414
HEUBATIODOSSC 779, 850 515, 077 343, 747 077, 319 85, 691 117, 093 82, 928 94, 494 HEUBATIODOSSC 73, 935 34, 803 80, 605 26, 562 28, 789 47, 944 60, 087 31, 519 HEUBATIODOSSC 73, 935 34, 866 35, 270 18, 733 6, 213 7, 458 7, 714 4, 624 HEUBATIODOST 50, 148 19, 281 37, 515 12, 222 19, 562 29, 874 22, 045 23, 135 HEUBATIODOST 50, 148 19, 281 37, 515 12, 222 19, 562 29, 874 22, 045 23, 135 HEUBATIODOST 50, 148 19, 281 37, 515 12, 222 19, 562 29, 874 22, 045 23, 135 HEUBATIODOST 41, 417 19, 132 24, 499 15, 544 4, 093 16, 474 41, 195 14, 145 HEUBATIODOST 90, 061 33, 532 39, 641 19, 754 27, 516 38, 801 31, 347 36, 440 HEUBATIODOSST 34, 417 19, 132 24, 499 19, 544 34, 385 38, 224, 897 29, 059 HEUBATIODOSST 79, 79, 79, 79, 79, 79, 79, 79, 79, 79,	15									
HEMBAT0009582		HEMBA1000946	15.461	5. 506	18.692	9.000				
HEMBAT0009582		HEMBA1000960	179 860	151.073	343.747	107.319	85, 691	117.093	82.928	94, 494
### HEBBA10009871								47 944		
HEBBA1000971 50,148 19,281 37,515 12,222 19,562 29,874 22,045 23,135 HEBBA1000974 51,100 33,405 188,137 28,372 24,575 23,751 13,731 27,272 HEBBA1000974 5,569 10,649 12,866 2,929 2,603 3,800 6,104 4,964 HEBBA1000975 34,417 19,132 42,499 15,644 4,009 18,478 14,192 14,353 HEBBA1000981 35,338 31,281 38,672 19,544 34,385 38,280 24,887 29,058 HEBBA1000981 35,338 31,281 38,672 19,544 34,385 38,280 24,887 29,058 HEBBA1000985 7,191 34,505 56,682 25,255 32,847 27,465 31,286 HEBBA1000986 72,707 55,780 160,717 34,676 32,494 41,317 23,483 37,846 HEBBA1000987 72,707 55,780 160,717 34,676 32,494 41,317 23,483 37,846 HEBBA1001007 12,569 42,553 50,717 34,676 32,494 41,317 23,483 37,846 HEBBA1001008 124,864 47,842 31,746 18,125 25,490 52,693 30,668 24,961 HEBBA1001009 37,843 29,289 33,565 23,255 34,263 47,777 4,955 48,154 HEBBA1001017 50,408 20,212 48,394 16,020 28,537 31,917 27,876 24,283 HEBBA1001017 50,408 20,212 48,394 16,020 28,537 31,917 27,876 24,283 HEBBA1001012 33,83 25,835 23,236 6,154 10,598 5,643 3,920 3,926 3,944 HEBBA1001012 33,83 25,835 23,236 6,154 10,598 5,643 3,920 3,926 3,926 HEBBA1001012 33,83 25,835 23,836 6,154 10,598 5,643 3,920 3,926 3,926 HEBBA1001012 33,83 25,835 23,836 8,343 8,013 3,917 27,876 24,283 HEBBA100102 33,83 25,835 23,836 8,357 33,556 33,337 3,946 6,154 10,598 5,643 3,920 3,926 3,936										
### PROPRIEST 10.0 33.45 18.17 17.27 28.97 24.576 23.715 13.731 27.27 ### PROPRIEST 10.0 33.46 18.171 28.97 24.576 23.715 13.731 27.27 ### PROPRIEST 10.0 33.46 19.28 18.171 28.97 24.576 23.715 13.731 27.27 ### PROPRIEST 19.132 42.499 15.644 4.009 16.478 41.192 14.353 ### PROPRIEST 19.132 42.499 15.644 4.009 16.478 41.192 14.353 ### PROPRIEST 19.132 42.499 15.644 4.009 16.478 41.192 14.353 ### PROPRIEST 19.132 42.499 15.644 4.009 16.478 41.192 14.353 ### PROPRIEST 19.132 42.499 15.644 4.009 16.478 41.192 14.353 ### PROPRIEST 19.132 42.499 15.642 27.151 38.801 31.347 36.440 ### PROPRIEST 19.132 39.641 19.754 27.151 38.801 31.47 36.440 ### PROPRIEST 19.132 39.641 19.754 27.515 38.801 31.47 36.440 ### PROPRIEST 19.132 39.641 19.554 27.515 38.801 31.47 36.440 ### PROPRIEST 19.132 39.641 19.554 27.515 39.641 39.015 7.652 31.286 ### PROPRIEST 19.132 39.641 39.6562 25.559 30.658 27.465 31.286 ### PROPRIEST 12.569 42.555 59.605 55.652 25.957 85.340 37.862 31.257 ### PROPRIEST 12.569 42.555 59.607 23.555 35.607 23.557 35.607 23.557		HEMBA1000968	14. 529	12. 486	35. 2/0					
HEMBAID00975 51, 100 33, 450 188, 137 28, 972 24, 576 23, 736 13, 731 27, 272 HEMBAID00975 34, 417 19, 132 42, 439 15, 644 4, 009 16, 478 14, 192 14, 353 HEMBAID00975 34, 417 19, 132 42, 439 15, 644 4, 009 16, 478 14, 192 14, 353 HEMBAID00978 30, 661 38, 512 39, 641 19, 754 27, 516 38, 261 31, 347 36, 440 HEMBAID00981 33, 318 31, 281 38, 672 19, 544 34, 185 38, 280 24, 897 29, 059 HEMBAID00985 73, 199 34, 501 58, 683 22, 640 32, 825 32, 184 27, 485 31, 286 HEMBAID00985 72, 707 55, 780 160, 717 40, 588 6, 343 90, 135 7, 852 3, 257 HEMBAID00985 72, 707 55, 780 160, 717 34, 676 32, 494 41, 317 23, 483 37, 846 HEMBAID001007 123, 590 42, 563 59, 807 23, 525 34, 283 47, 777 47, 495 48, 154 HEMBAID001008 124, 864 47, 842 83, 746 18, 125 25, 490 52, 693 30, 668 24, 961 HEMBAID001008 124, 864 47, 842 83, 746 18, 125 25, 490 52, 693 30, 668 24, 961 HEMBAID001009 73, 737 7, 582 14, 865 61, 54 10, 598 5, 643 3, 920 7, 188 HEMBAID001010 109, 049 83, 356 233, 234 60, 123 61, 977 94, 424 47, 095 74, 625 HEMBAID00101 109, 049 83, 356 233, 234 60, 123 61, 977 94, 424 47, 095 74, 625 HEMBAID00102 53, 667 55, 646 31, 540 25, 647 24, 596 23, 146 27, 169 HEMBAID00102 53, 667 55, 646 31, 540 25, 647 24, 596 23, 146 27, 169 HEMBAID00102 53, 657 55, 646 11, 816 31, 840 25, 647 24, 596 24, 183 HEMBAID00102 37, 833 29, 29, 20 17, 500 19, 183 9, 299 8, 047 37, 20 HEMBAID00102 37, 833 25, 835 28, 969 18, 452 20, 770 22, 790 25, 194 20, 783 HEMBAID00102 37, 838 25, 835 28, 969 18, 540 25, 647 24, 596 24, 184 HEMBAID00102 37, 838 25, 835 28, 969 18, 184 26, 22, 197 24, 250 HEMBAID00103 30, 248 24, 251 31, 348 31, 860 33, 390 34,			50, 148	19, 281	37, 515	12, 222	19, 562	29. 874	22.045	23. 135
HEMBA1000975 34.417 19.312 42.499 15.644 4.009 16.478 14.192 14.353 HEMBA1000979 30.061 38.532 99.641 19.754 27.516 38.801 31.347 36.440 HEMBA1009983 71.391 34.501 58.672 79.544 34.185 38.280 42.897 29.059 HEMBA100988 71.991 34.501 58.683 22.640 79.2825 32.384 27.485 31.286 HEMBA1000988 71.991 34.501 58.683 22.640 79.2825 32.384 27.485 31.286 HEMBA1000988 79.290 20.363 22.497 4.058 6.343 9.035 7.852 3.257 HEMBA1000989 72.707 55.780 150.717 34.676 32.494 41.317 23.483 37.846 HEMBA1001007 123.690 42.563 99.807 74.525 32.257 HEMBA1001007 123.690 42.563 99.807 74.525 32.257 HEMBA1001007 123.690 42.563 99.807 74.525 34.283 47.777 47.496 48.154 HEMBA1001007 123.690 42.563 99.807 72.525 34.283 47.777 47.496 48.154 HEMBA1001007 123.690 42.563 99.807 25.255 34.283 47.777 47.496 48.154 HEMBA1001007 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEMBA1001010 90.409 83.356 231.234 60.123 61.977 94.424 47.095 74.625 HEMBA1001010 97.327 7.582 48.394 60.123 61.977 94.424 47.095 74.625 HEMBA1001010 97.327 7.582 48.394 60.123 61.977 94.424 47.095 74.625 HEMBA1001010 97.327 7.582 48.394 60.123 61.977 94.424 47.095 74.625 HEMBA1001020 53.687 55.646 15.814 30.598 5.643 3.920 7.188 HEMBA1001021 115.724 42.415 59.434 60.123 61.977 94.424 47.095 74.625 14.865 HEMBA1001020 53.687 55.646 115.814 31.640 25.647 24.196 23.146 27.169 HEMBA1001022 115.724 42.415 59.434 28.28 28.1818 64.484 64.173 29.614 HEMBA1001022 33.524 15.383 25.835 28.998 18.452 20.270 22.970 25.194 20.783 HEMBA1001022 33.524 15.335 16.511 8.023 11.818 13.894 8.606 8.998 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.998 HEMBA1001025 33.899 13.800 19.077 12.856 28.282 20.270 22.970 25.194 20.783 HEMBA1001075 33.254 18.800 19.077 12.855 18.851 18.452 44.382 15.726 38.599 18.452 18.452 48.859 18.599 18.452 18.452 95.500 24.452 56.356 68.994 HEMBA1001076 33.254 18.350 18.350 18.351 1										
HEBBA1000975 34.417 19.12 42.499 15.544 4.009 16.478 14.192 14.353 HEBBA1000975 99.061 38.532 99.641 19.754 27.516 38.801 31.347 36.440 HEBBA1000981 33.338 31.381 33.672 19.544 34.385 38.200 24.897 29.059 HEBBA1000983 71.391 34.501 58.683 22.640 32.825 32.384 27.485 31.286 HEBBA1000986 79.290 20.353 22.497 4.058 6.343 9.035 7.582 3.257 HEBBA1000986 79.290 70.353 22.497 4.058 6.343 9.035 7.582 3.257 HEBBA1000986 79.290 70.557.80 160.717 34.676 32.494 41.317 23.83 37.846 HEBBA1000986 79.27 77 55.780 160.717 34.676 32.494 41.317 23.83 37.846 HEBBA1001009 72.707 55.780 160.717 34.676 32.494 41.317 23.83 37.846 HEBBA1001009 37.843 29.269 33.746 18.125 25.490 52.693 30.668 24.961 HEBBA1001009 37.843 29.269 35.6715 11.055 17.115 17.937 17.701 22.055 HEBBA10010101 109.049 83.356 233.526 31.575 17.115 17.937 17.701 22.055 HEBBA1001010 109.049 83.356 233.526 31.60123 60.123	20									
HEMBA1000981 33. 318 31. 281 38. 672 19. 544 4. 009 16. 478 14. 192 14. 353 HEMBA1000981 35. 338 31. 281 38. 672 19. 544 34. 385 38. 280 24. 897 29. 059 HEMBA1000985 71. 391 34. 501 58. 683 22. 640 72. 825 32. 384 27. 685 31. 286 HEMBA1000986 72. 71. 391 34. 501 58. 683 22. 640 72. 825 32. 384 27. 685 31. 286 HEMBA1000986 72. 71. 79 57. 680 56. 717 34. 676 57. 85. 340 63. 718 54. 6892 HEMBA1000981 72. 707 55. 780 56. 717 34. 676 52. 957 85. 340 63. 718 54. 6892 HEMBA1001009 72. 707 55. 780 56. 717 34. 676 52. 957 85. 340 63. 718 54. 6892 HEMBA1001008 124. 864 47. 842 83. 746 18. 125 25. 490 52. 583 30. 668 24. 956 HEMBA1001008 37. 843 29. 269 35. 715 11. 055 71. 715 71. 737 74. 74. 96 48. 154 HEMBA1001010 109. 049 83. 355 231. 234 60. 123 61. 977 94. 424 47. 095 74. 625 HEMBA100101 109. 049 83. 355 231. 234 60. 123 61. 977 94. 424 47. 095 74. 625 HEMBA1001019 7. 127 7. 582 14. 865 61. 54 10. 598 5. 643 3. 920 71. 88 HEMBA1001022 37. 883 32. 835 53. 836 38. 678 25. 647 02. 25. 597 27. 876 64. 71. 699 HEMBA1001022 37. 883 32. 835 53. 836 38. 678 25. 647 02. 270 22. 790 25. 194 20. 733 HEMBA1001022 37. 883 32. 835 53. 835 38. 628 25. 181 64. 484 64. 173 29. 814 HEMBA1001022 37. 883 32. 835 38. 836 18. 628 25. 181 64. 484 64. 173 29. 814 HEMBA1001022 37. 883 32. 835 53. 836 18. 628 25. 181 64. 646 64. 173 29. 814 HEMBA1001022 37. 883 32. 835 53. 835 18. 638 7. 288 12. 653 8419 7. 418 HEMBA1001023 37. 838 32. 835 53. 837 32. 838 7. 288 12. 653 8. 419 7. 418 HEMBA1001024 37. 883 32. 835 37. 846 18. 125 38. 849 38. 856 38. 819 38. 856 38. 819 38. 856 38. 819 38. 819 38. 856 38. 819 38. 856 38. 819 39. 89 39.	-	HEMBA1000974	5.609	10.649		2.929				
HEMBAIDOOD981 35, 338 31, 281 38, 672 39, 641 19, 754 27, 516 38, 801 31, 347 36, 440			34 417	19, 132	42 499	15.644	4, 009	16.478	14. 192	14.353
HEMBA1000981 35. 338 31. 281 38. 672 39. 544 34. 385 38. 280 24. 897 29. 059 HEMBA1000983 71. 391 34. 501 58. 683 22. 640 32. 825 32. 384 27. 465 31. 286 HEMBA1000986 72. 714 74. 713 236. 019 56. 662 52. 957 85. 340 63. 718 54. 892 HEMBA1000996 72. 707 55. 780 160. 717 34. 676 32. 494 41. 317 23. 483 37. 846 HEMBA1001007 72. 690 42. 563 69. 807 23. 525 34. 263 47. 777 47. 495 48. 154 HEMBA1001008 72. 869 42. 563 69. 807 23. 525 34. 263 47. 777 47. 495 48. 154 HEMBA1001008 72. 864 47. 842 33. 746 18. 125 25. 490 52. 693 30. 685 43. 961 HEMBA1001008 73. 843 29. 269 36. 715 11. 055 71. 115 77. 937 77. 701 22. 055 HEMBA1001014 109. 049 83. 356 233. 234 60. 123 61. 977 94. 424 47. 095 74. 625 HEMBA1001017 50. 408 20. 212 48. 394 16. 202 28. 537 31. 917 27. 767 24. 283 HEMBA1001019 7. 327 7. 582 14. 865 6. 154 10. 598 5. 643 3. 920 7. 188 HEMBA1001022 33. 687 35. 686 15. 818 31. 640 25. 647 24. 596 23. 146 27. 169 HEMBA1001022 37. 883 25. 835 28. 969 18. 452 20. 270 22. 790 25. 194 20. 783 HEMBA1001022 37. 883 25. 835 28. 969 18. 452 20. 270 22. 790 25. 194 20. 783 HEMBA1001022 33. 892 13. 850 19. 667 12. 855 11. 485 9. 299 8. 047 8. 898 HEMBA1001022 38. 892 13. 850 19. 667 12. 855 11. 445 24. 382 15. 726 12. 323 HEMBA1001052 38. 893 13. 850 19. 667 12. 855 11. 485 31. 894 8. 806 8. 998 HEMBA1001052 38. 892 74. 625 56. 555 72. 625 72. 625 73. 38. 575 52. 512 HEMBA1001077 24. 869 115. 813 33. 143 100. 376 67. 510 13. 193 67. 92. 97 38. 575 52. 512 HEMBA1001078 33. 254 26. 761 41. 713 26. 498 24. 531 48. 419 7. 418 HEMBA1001078 33. 254 26. 761 41. 713 26. 498 24. 531 48. 44 47. 276 36. 510 HEMBA1001079 30										
HEMBAID00983										
		HEMBA1000981	35.338	31. 281	38.672				24.897	
HEMBAI000985 9,290 20,363 22,497 4,058 5,343 9,015 7,852 3,257 HEMBAI000986 128,714 74,713 236,019 56,662 52,957 85,340 63,718 54,852 HEMBAI000999 72,707 55,780 160,717 34,676 32,494 41,317 23,483 37,846 HEMBAI001007 123,690 42,563 69,807 23,525 34,263 47,777 47,496 48,154 48,164 48,154 48,164		HEMRATOGOGS		34 501	58 683	22,640	32.825	32, 384	27.465	31, 286
HEMBAI001098 128, 714 74, 713 236, 019 56, 662 55, 957 85, 340 63, 718 54, 832 HEMBAI001099 72, 707 75, 780 160, 717 34, 676 32, 494 41, 317 23, 483 37, 846 HEMBAI001008 124, 864 47, 842 83, 746 18, 125 25, 490 52, 693 30, 668 24, 961 HEMBAI001009 37, 843 29, 269 36, 715 11, 055 77, 157 79, 701 77, 701 22, 055 HEMBAI001014 109, 049 83, 356 213, 234 60, 123 61, 977 94, 424 47, 095 74, 625 74,										
HEMBAI000986 128.714 74.713 238.019 38.002 32.314 31.00 63.718 34.02 34.00 63.718 34.002 34.000 37.845 34.000 37.845 34.000 37.845 34.000 37.845 34.000 37.845 34.000 37.845 34.000 37.843 37.845 34.000 37.843 37.845 34.000 37.843 37.845 34.000 37.845 34.000 37.843 39.000 37.845 39.000 39.00	25									
HEMBAIODIOO7 123.680 42.563 69.807 23.525 34.263 47.777 47.496 48.154 HEMBAIODIOO8 124.864 47.842 83.746 18.125 25.490 52.693 30.668 24.961 HEMBAIODIOO9 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEMBAIODIO14 109.049 83.356 231.234 60.123 61.977 94.424 47.095 74.625 HEMBAIODIO17 50.408 20.212 48.394 16.020 28.537 31.917 27.876 24.283 HEMBAIODIO19 7.327 7.582 14.865 6.154 10.598 5.643 3.920 7.188 HEMBAIODIO109 7.327 7.582 14.865 6.154 10.598 5.643 3.920 7.188 HEMBAIODIO20 53.067 55.646 115.814 31.640 25.647 24.996 23.146 27.169 HEMBAIODIO22 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBAIODIO22 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBAIODIO33 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBAIODIO51 124.869 115.181 387.345 100.376 67.510 19.163 9.299 8.047 8.589 HEMBAIODIO59 98.097 41.525 66.565 27.826 26.220 46.725 42.366 36.506 HEMBAIODIO69 38.097 41.525 66.565 27.826 26.220 46.725 42.366 36.506 HEMBAIODIO69 32.251 24.026 33.337 14.007 7.907 13.710 17.387 15.720 HEMBAIODIO69 33.254 25.761 41.4713 26.498 24.531 31.498 25.302 23.636 HEMBAIODIO69 32.251 24.026 33.337 14.007 7.907 13.710 17.387 15.720 HEMBAIODIO69 32.251 24.026 33.337 14.007 7.907 13.710 17.387 15.720 HEMBAIODIO69 59.8097 41.525 66.565 27.826 26.220 46.725 42.366 36.506 HEMBAIODIO69 57.701 23.951 31.254 22.489 24.848 33.265 31.880 12.873 HEMBAIODIO77 24.689 23.055 64.486 19.413 16.821 16.821 16.858 13.165 12.873 HEMBAIODIO68 57.701 23.951 31.254 22.899 24.848 33.265 31.880 12.873 HEMBAIODIO79 33.3254 25.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBAIODIO79 36.838 41.625 17.1096 31.438 31.760 24.829 17.487 25.551 12.873 12.874 18.884 18.000 1006 57.701 29.951 31.254 22.899 24.848 33.265 31.880 26.484 18.800 1006 57.701 29.951 31.254 22.899 24.488 33.265 31.880 26.484 18.800 1006 57.701 29.951 31.265 19.406 24.829 17.487 25.551 18.507 18.500 19.600 19.600 19.500	23	HEMBA1000986	128.714	74. 713	236.019	56.662	52.957	85. 340	63.718	54.892 -
HEMBAIODIOO7 123.680 42.563 69.807 23.525 34.263 47.777 47.496 48.154 HEMBAIODIOO8 124.864 47.842 83.746 18.125 25.490 52.693 30.668 24.961 HEMBAIODIOO9 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEMBAIODIO14 109.049 83.356 231.234 60.123 61.977 94.424 47.095 74.625 HEMBAIODIO17 50.408 20.212 48.394 16.020 28.537 31.917 27.876 24.283 HEMBAIODIO19 7.327 7.582 14.865 6.154 10.598 5.643 3.920 7.188 HEMBAIODIO109 7.327 7.582 14.865 6.154 10.598 5.643 3.920 7.188 HEMBAIODIO20 53.067 55.646 115.814 31.640 25.647 24.996 23.146 27.169 HEMBAIODIO22 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBAIODIO22 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBAIODIO24 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBAIODIO33 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBAIODIO51 124.869 115.181 387.345 100.376 67.510 19.163 9.299 8.047 8.589 HEMBAIODIO59 98.097 41.525 66.565 27.826 26.220 46.725 42.366 36.506 HEMBAIODIO69 38.097 41.525 66.565 27.826 26.220 46.725 42.366 36.506 HEMBAIODIO69 32.251 24.026 33.337 14.007 7.907 13.710 17.387 15.720 HEMBAIODIO69 33.254 25.761 41.4713 26.498 24.531 31.498 25.302 23.636 HEMBAIODIO69 32.251 24.026 33.337 14.007 7.907 13.710 17.387 15.720 HEMBAIODIO69 32.251 24.026 33.337 14.007 7.907 13.710 17.387 15.720 HEMBAIODIO69 59.8097 41.525 66.565 27.826 26.220 46.725 42.366 36.506 HEMBAIODIO69 57.701 23.951 31.254 22.489 24.848 33.265 31.880 12.873 HEMBAIODIO77 24.689 23.055 64.486 19.413 16.821 16.821 16.858 13.165 12.873 HEMBAIODIO68 57.701 23.951 31.254 22.899 24.848 33.265 31.880 12.873 HEMBAIODIO79 33.3254 25.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBAIODIO79 36.838 41.625 17.1096 31.438 31.760 24.829 17.487 25.551 12.873 12.874 18.884 18.000 1006 57.701 29.951 31.254 22.899 24.848 33.265 31.880 26.484 18.800 1006 57.701 29.951 31.254 22.899 24.488 33.265 31.880 26.484 18.800 1006 57.701 29.951 31.265 19.406 24.829 17.487 25.551 18.507 18.500 19.600 19.600 19.500		HEMBAINING 1	72 707	55 780	160 717	34 676	32, 494	41.317	23, 483	37.846
HEMBA1001008 124.864 47.842 83.746 18.125 25.490 52.693 30.668 24.967 HEMBA1001009 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEMBA1001014 109.049 83.356 233.234 60.123 61.977 94.424 47.095 74.625 HEMBA1001017 50.408 20.212 48.394 16.020 28.537 31.917 27.876 24.283 HEMBA1001019 7.327 7.582 14.865 6.154 10.598 5.643 31.917 27.876 24.283 HEMBA1001020 53.067 55.646 115.814 31.540 25.647 24.598 23.146 27.169 HEMBA1001021 115.724 42.415 59.434 28.828 26.181 64.484 64.173 29.614 HEMBA1001022 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001026 21.343 12.515 18.851 6.888 7.288 12.663 8.419 7.418 HEMBA1001025 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001055 38.097 41.525 66.565 27.826 26.220 46.725 42.566 36.506 HEMBA1001066 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.512 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001067 23.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001066 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.512 HEMBA1001060 57 701 23.951 31.254 24.898 24.531 31.498 25.302 23.636 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 25.581 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 25.581 HEMBA1001098 57.701 23.951 31.254 22.489 24.531 31.498 25.302 23.636 HEMBA1001099 18.322 24.021 44.537 46.695 19.266 25.446 4.525 31.098 44.625 31.099 31.438 31.760 24.829 17.487 25.581 HEMBA1001099 18.322 24.021 48.14 7.146 13.778 16.655 17.044 10.190 HEMBA1001099 18.332 22.40 24.42,547 77.235 81.846 4.555 31.468 33.498 44.927 25.531 HEMBA1001099 18.322 24.021 48.814 7.146 13.450 93.324 78.590 18.522 11.099 18.722 24.021 48.814 7.146 13.778 16.655 17.044 10.190 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559										
HEMBA1001009 37.843 29.269 36.715 11.055 17.115 17.937 17.701 22.055 HEMBA1001014 109.049 83.356 233.234 60.123 61.977 94.424 47.095 74.625 17.626 19.000 19.049 83.356 233.234 60.123 61.977 94.424 47.095 74.625 17.626 19.000 19.049 83.356 233.234 60.123 61.977 94.424 47.095 74.625 17.626 19.000										
HEMBA1001009 37. 843 29. 269 35. 715 11. 055 17. 115 17. 937 17. 701 22. 055 HEMBA1001014 109. 049 83. 356 233. 234 60. 123 61. 977 94. 424 47. 095 74. 625 HEMBA1001017 50. 408 20. 212 48. 394 16. 020 28. 537 31. 917 27. 876 24. 283 HEMBA1001019 7. 327 7. 582 14. 865 6. 154 10. 598 5. 643 3. 920 7. 188 HEMBA1001020 53. 067 55. 646 115. 814 31. 640 25. 647 24. 596 23. 146 27. 169 HEMBA1001022 37. 883 25. 835 28. 969 18. 452 20. 270 22. 790 25. 194 20. 783 HEMBA1001024 23. 524 15. 235 16. 511 8. 023 11. 818 13. 894 8. 606 8. 098 HEMBA1001024 23. 524 15. 235 16. 511 8. 023 11. 818 13. 894 8. 606 8. 098 HEMBA1001043 10. 374 11. 995 9. 892 10. 750 19. 163 9. 299 8. 047 8. 589 HEMBA1001051 124. 869 115. 181 387. 345 100. 376 67. 510 61. 660 46. 295 68. 994 HEMBA1001053 38. 992 13. 860 19. 067 12. 855 11. 445 24. 382 15. 726 23. 23 HEMBA1001059 38. 997 41. 525 66. 565 27. 826 26. 220 46. 725 42. 356 36. 506 HEMBA1001060 116. 857 74. 020 161. 485 61. 750 50. 524 52. 957 38. 575 52. 612 HEMBA1001077 25. 850 16. 043 19. 924 7. 855 3. 425 9. 530 6. 779 24. 242 HEMBA1001077 24. 689 23. 055 64. 486 19. 413 16. 821 16. 858 13. 165 12. 873 HEMBA1001080 57. 701 23. 951 31. 254 22. 489 24. 848 33. 265 31. 880 26. 484 HEMBA1001081 57. 520 16. 043 19. 924 7. 855 3. 425 9. 530 6. 779 24. 242 HEMBA1001080 57. 701 23. 951 31. 254 22. 489 24. 848 33. 265 31. 880 26. 484 HEMBA1001081 57. 520 16. 043 19. 924 7. 855 3. 425 9. 530 6. 779 24. 242 HEMBA1001080 57. 701 23. 951 31. 254 22. 489 24. 848 33. 265 31. 880 26. 484 HEMBA1001081 59. 252 116. 909 94. 247 77. 235 81. 386 44. 521 48. 34 HEMBA1001081 59. 252 16. 909 94		HEMBA1001008	124.864	47.842	83.746	18.125	25.490	52.693	30.668	24.961
HEMBA1001014			37 843			11 055	17 115	17, 937	17,701	22.055
HEMBA1001017										
HEMBA1001019	20									
HEMBA1001019	30	HEMBA1001017	50.408	20. 212	48. 394	16.020	28.537	31.917	27.876	24. 283
#EMBA1001020					14 865	6 154	10 598	5, 643	3, 920	7 188
HEMBA1001021 115.724 42.415 59.434 28.828 26.181 64.484 64.173 29.614 HEMBA1001022 37.883 25.835 28.969 18.452 20.270 22.790 25.194 20.783 HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001026 21.343 12.515 18.851 5.888 7.288 12.663 8.419 7.418 HEMBA1001043 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBA1001052 138.892 115.818 1387.345 100.376 67.510 61.660 46.295 68.994 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001060 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001085 33.254 26.761 41.713 26.498 24.531 31.498 25.302 23.636 HEMBA1001086 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001085 77.01 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001085 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.531 HEMBA1001085 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.531 HEMBA1001089 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001093 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001093 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001093 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001109 186.384 190.240 540.908 155.496 134.530 93.324 78.690 1116.187 HEMBA10011121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 32.00 65.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001123 108.859 55.807 190.785 414 145 39.028 42.683 25.551 30.174 HEMBA1001123 108.859 55.807 190.785 414 145 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 3										
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HEMBA1001024 23.524 15.235 16.511 8.023 11.818 13.894 8.606 8.098 HEMBA1001026 21.343 12.515 18.851 5.838 7.288 12.663 8.419 7.418 HEMBA1001043 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBA1001051 124.869 115.181 387.345 100.376 67.510 61.660 46.295 68.994 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001060 116.857 74.020 161.485 61.750 50.524 52.957 38.575 52.612 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001071 25.850 16.043 19.924 7.855 3.425 9.530 6.779 24.242 HEMBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001078 33.254 26.761 41.713 25.498 24.531 31.498 25.302 23.636 HEMBA1001080 57.701 23.951 31.254 22.489 24.848 33.265 31.880 26.484 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001093 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA10011094 5.535 8.779 10.059 3.089 4.628 4.521 4.834 4.468 HEMBA10011094 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.883 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367					28 969	18 452	20 270	22 790	25 194	20 783
HEMBA1001026 21.343 12.515 18.851 6.888 7.288 12.663 8.419 7.418 HEMBA1001043 10.374 11.995 9.892 10.750 19.163 9.299 8.047 8.589 HEMBA1001051 124.869 115.181 387.345 100.376 67.510 61.660 46.295 68.994 HEMBA1001052 38.892 13.860 19.067 12.855 11.445 24.382 15.726 12.323 HEMBA1001059 98.097 41.525 66.565 27.826 26.220 46.725 42.356 36.506 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001064 32.251 24.026 33.937 14.007 7.907 13.710 17.387 16.720 HEMBA1001077 24.689 23.055 64.486 19.413 16.821 16.858 13.165 12.873 HEMBA1001084 62.698 41.625 171.096 31.438 31.760 24.829 17.487 26.581 HEMBA1001085 159.252 116.909 294.247 77.235 81.384 76.498 59.989 55.574 HEMBA1001088 74.704 42.537 46.695 19.266 25.146 33.498 44.927 26.310 HEMBA1001093 30.048 28.810 72.081 20.831 14.610 11.033 15.558 22.531 HEMBA1001094 5.535 8.779 10.059 3.083 4.628 4.521 4.834 4.468 HEMBA1001099 18.322 24.021 14.814 7.146 13.778 16.055 11.044 10.190 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 21.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 27.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 27.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 27.559 18.527 HEMBA1001104 21.919 13.788 35.048 9.637 18.058 24.450 27.559 18.527 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
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HEMBA1001114 89.023 252.529 187.547 75.887 35.109 66.259 69.432 341.702 HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367	50	HEMBA1001109	į 186. 384	190, 240	540.908	155. 496	134.530		1 /8. 590	116.187
HEMBA1001121 32.820 25.812 89.860 19.710 34.244 18.209 9.519 15.621 HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367				252 529	187 547	75, 857	35 109	66.259	69.432	341, 702
HEMBA1001122 3.304 6.213 8.316 4.763 19.120 5.650 4.506 23.059 HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367										
HEMBA1001123 108.859 55.807 190.789 41.415 39.028 42.683 25.551 30.174 HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367		HEMBA1001122	3. 304	6.213	8.316	4.763	19.120	5. 650	4.506	¿ 23.059
HEMBA1001133 50.744 21.167 36.786 14.764 34.752 26.702 23.524 11.367		HEMMATON 199			190 789			42.683	25.551	30, 174
		DEMONSTRATES								
55 HEMBATOOT137 38.685 21.659 46.297 21.567 13.174 15.867 11.767 25.508										
	55		38. 685	21.659	46.297	21.567	13.174	15.867	11.767	25. 508
										· · · · · · · · · · · · · · · · · · ·

Table 17

	DE USA 100 LA	- A	56 168 1		40 037	60 363 1	44 463 1	20 207	12 551
	HEMBA1001140	60.453	66.122	169.353	48.837	60.363	44. 403	30.367	43.561
	HEMBA1001144	278.126	195.811	643.688	207. 291	166.089	101.134	106.337.	142. 120
5	HEMBA1001145	58.539	241.368	206.084	46. 342	39. 316	61.827	91.170	66.852
5	HEMBA1001158	29.417	28. 121	43.877	13.337	24.176	19.965	18.089	28, 622
	HEMBA1001172	74.727	47.695	213.708	37.115	24. 460	26, 620	19.178	32.709
	HEMBA1001174				7.914	2. 574	8.031	3.119	4. 980
		6. 279	8.617	8.831					
	HEMBA1001175	29.561	34.909	43. 568	19.819	34. 829	16. 588	19.883	17. 824
	HEMBA1001182	136.762	64.608	105.979	44.066	83.417	86.736	126. 297	79. 785
10	HEMBA1001184	16.758	9,703	22.060	9.016	11.018	10.205	6.347	9. 176
10	HEMBA1001192	15.119	10.798	11.626	6.559	5. 736	3. 435	9.089	11.273
	HEMBA1001197	82.571	114.743	110.687	83. 431	56.396	68.797	99.959	173, 379
	HEMBA1001208	40.250	30, 964	37.220	19. 514	11.451	24. 172	27.637	12. 469
								51.978	
	HEMBA1001213	81.501	37. 345	57.618	18.958	24.480	52. 160		31. 326
	HEMBA1001214	36.798	16,011	20.958	17.612	12.418	20.697	19.108	21. 328
15	HEMBA1001221	14.108	10. 456	11.382	7.001	17.058	10.307	7. 980	11. 111
,,	HEMBA1001225	13.961	14.077	13.384	5.925	5. 876	13.456	12.076	5. 825
	HEMBA1001226	173.501	137.685	444.754	120.060	113.306	75. 167	63.960	67. 304
	HEMBA1001228	115.971	48,677	102.518	36.755	64.214	50.002	60.915	35, 178
	HEMBA1001229	246.802	111.161,	135.886	43.460	94.703	148.387	156.871	115. 302
	HEMBA1001235	43.880	86.102	81.818	36.769	54. 172	65.830	70.065	66. 201
20	HEMBA1001238	67. 342	62.561	136. 273	36.471	33.652	41.838	26.195	28.747
	HEMBA1001242	55. 562	43. 106	58. 593	41. 382	47. 200	38.498	43.114	44. 230
	HEMBA1001247	28.768	22. 129	16.518	10.576	8.758	17.031	9.651	13. 385
	HEMBA1001253	58. 130	60.415	66.640	18. 982	45. 992	54.071	95.073	63.393
	HEMBA1001257	33.557	18.509	24. 256	10.657	12.732	31.261	24.849	9.134
	HEMBA1001261	585.214	143.415	243.791	98. 186	169. 988	310.109	234.388	125. 796
	HEMBA1001262	27.336	17.339	19.088	5.647	15.678	20.899	11.464	19.889
25	HEMBA1001265	36.604	28.090	152.221	27, 730	49, 893	34. 423	16.502	26.993 -
	HEMBA1001266	69.367	67.414	170.657	45. 898	31.802	39, 554	41.287	52, 480
	HEMBA1001269	69.921	44.649	36.964	34, 126	22. 232	42.207	49.848	39.719
	HEMBA1001272	20. 406	15.416	11.514	7. 843	8. 604	7.893	20.960	13. 545
				147. 371	41.113	50.841	58. 248	43. 344	47.548
	HEMBA1001279	113.597	76.085						
30	HEMBA1001281	45. 326	37.551	65. 225	44. 536	46.787	41.371	32.229	56.625
30	HEMBA1001286	370.697	150.949	236.623	103. 571	123.976	219, 461	196.233	117. 566
	HEMBA1001289	41.041	24.670	40.151	15. 175	30.612	27.627	26.637	19.344
	HEMBA1001291	76.537	40.444	50. 226	18.775	38. 423	55.355	46.692	35. 972
	HEMBA1001294	82.258	72.319	157.642	42.143	20. 735	29. 333	17.711	34. 443
	HEMBA1001296	53.487	17.150	31.045	10. 275	15.918	21, 120	15.842	13. 595
	HEMBA1001297	13.397	24.306	19.513	11.631	14.701	4. 543	9.800	8. 121
35	HEMBA1001259	122.378	135, 140	326.747	90.817	73.749	56.152	49.803	80.999
	HEMBA1001302	56.839	29.036	56.412	19.108	20.078	34. 481	51.929	37.087
	HEMBA1001303	14. 975	18.442	43.778	16.797	10.985	11.442	9,787	19.264
	HEMBA1001306	262.869	135.864	244. 234	109.949	109. 582	147.334	146.509	115, 543
	HEMBA1001308	174.017	96. 705	220.049	56. 953	61, 486	74. 225	56, 171	58.657
	HEMBA1001310		52.915	67.714	22.895	38. 245	67. 233	49. 204	51,006
4.5		103.029		61.080	18.118	33, 555	47.007	41.795	38. 527
40	HEMBA1001312	98.664	47. 333				3.003	5. 682	3.780
	HEMBA1001319	2. 396	8.234	13.960	1.828	5. 485			
	HEMBA1001322	139.794	39.912	105.709	27.700	41. 977	70.428	70.602	46. 470
	HEMBA1001323	33.347	16.728	25. 356	11. 399	17. 982	11.181	6, 356	12.033
	HEMBA1001326	86.190	37. 984	69. 933	24. 331	30.078	49. 223	46.365	16.347
	HEMBA1001327	7. 232	9. 387	23.180	7.314	5. 185	9. 563	4.423	5. 267
45	HEMBA1001330	115.768	106.951	275.315	73. 389	24.661	70.535	40.088	77.680
•	HEMBA1001348	15.770	21.874	26.347	9. 575	13.666	23.703	12.647	13.724
	HEMBA1001350	75. 857	38.749	51.454	16.428	34, 291	56.400	34.055	24.753
	HEMBA1001351	52. 274	55.313	56.544	30. 521	46, 408	29.604	44.212	30. 972
	HEMBA1001352	68. 321	46.617	54. 427	17.559	29. 887	39.484	52.789	29. 131
	HEMBA1001353	39.891	57. 492	54.971	31. 425	27.945	45. 687	29.741	66. 188
				59.774			47.618	32.581	59. 101
50	HEMBA1001358	45.659	52.406		46.865	40. 225		13. 938	
	HEMBA1001361	22.908	16.519	28.635		15.569	13.635		16. 914
	HEMBA1001364	18.896	17. 205	23. 355		9.469	13.379	76. 125	15. 026
	HEMBA1001375	61.506	22. 179	38.795		25.778	40.077	21.715	22. 300
	HEMBA1001377	140. 430	131.029	307.084	83. 191	100.026	74.475	63.988	96. 351
	HEMBA1001383	23.974	26. 206	28.704	11.442	17.819	19.160	16.899	7.766
55	HEMBA1001387	58. 343	34. 130	63.677	19. 556	30.371	42. 397	40. 247	49. 239
ua									

Table 18

	FUENDA LAGRAGA T			20 077 [10 0co I	10 (24)	20 022 1	22 224	33.218
	HEMBA1001388	48.601	24.690	39. 877	18. 958	10.634	32.922	22. 224	57. 372
	HEMBA1001390	132.003		254. 352	56.412	64. 490	47. 169	44. 169	5. 544
5	HEMBA1001391	18. 302	9.686	12. 994	6. 299	10.600	8. 500	7.116	29, 984
	HEM8A1001398	91.232	50. 992	142.408	36.081	29. 548	29. 490	28.704	
	HEMBA1001405	58.645	22. 354	32. 227	15.864	9. 285	19, 993	24. 564	13.964
	HEMBA1001406	36.434	22.693	105. 808	18.094	19. 994	13.316	18.019	16. 592
	HEMBA1001407	38. 781	19.637	24. 599	18.935	13. 107	23.014	18. 826	15.060
	HEM8A1001411	28. 412	7.180	21.950	8. 303	9. 708	14. 302	8. 598	6.663
10	HEMBA1001413	66.736	26. 480	35. 635	15. 400	24.013	18. 356	24. 304	20.759
	HEMBA1001414	20.720	7.567	18. 414	12. 522	9. 722	12.903 32.235	18. 283	18. 581
	HEMBA1001415	76.802	54. 702	159.510	34. 156	20. 989	24. 993	21.694	26. 676
	HEMBA1001416	41.784	23.474	29. 453	12. 230	24. 881		25. 847 66. 751	28.651
	HEMBA1001432	74.066	60.077	190.870	40. 409	63.619	36.879 50.447	37.821	33. 675 64. 403
	HEMBA1001433	132.672	110. 163	246. 542	77.852	61.676	59.723	58. 393	56. 483
15	HEMBA1001435	138.669	108.645	334. 104	89. 523	15.704	7. 291	6.742	6. 336
	HEMBA1001442	13. 093	8. 504	11. 177	7. 985	27.976	37. 353	30. 266	41.647
	HEMBA1001446	102. 450	63. 255	146, 442	40. 086 30. 799	31. 322	42. 457	42.764	41. 349
	HEMBA1001450	72.339	35. 494	55. 103	88.679	43. 129	54, 712	41.131	31. 250
	HEMBA1001454	146.726	128.060 e 8.197	438. 247 8. 325	5. 561	4. 437	5. 252	4. 300	7. 359
	HEMBA1001455	5. 879 17. 432	15, 927	16, 490	6.749	2.733	5, 888	7.836	10.963
20	HEMBA1001459 HEMBA1001461	61,531	52, 734	57. 136	38. 874	24.764	19. 473	23. 241	32.318
	HEMBA1001462	10, 875	14, 911	16, 843	12. 984	13.465	48, 381	7.061	25. 992
	HEMBA1001463	137. 907	83. 753	340. 496	93.114	51.866	61.784	37.705	68. 960
	HEMBA1001469	85.416	21. 757	29, 463	15. 911	84, 887	77, 440	27.033	29. 537
	HEMBA1001473	20. 582	31.855	36.498	8. 307	3, 680	16.703	21.371	19.890
	HEMBA1001476	135, 720	113.851	246, 800	65, 595	57, 431	63.903	65. 229	67.697
25	HEMBA1001477	5. 228	2,001	4.505	2.645	1,540	3. 243	1.426	2.876
	HEMBA1001478	14, 335	10, 180	12.692	5. 468	4.474	5. 444	2. 171	4. 539
	HEMBA1001480	88.891	28, 381	49.689	21.660	14. 126	36. 334	38. 272	30. 563
	HEMBA1001483	29.872	5. 156	20. 900	4. 547	5, 264	9. 545	13.805	4. 424
	HEM8A1001490	6.867	6. 967	14. 148	7.289	1. 585	5.016	5. 792	5. 999
30	HEMBA1001495	431.282	118.073	203.714	73. 985	176.836	195. 947	194. 164	146. 945
30	HEMBA1001497	93.817	60. 807	227. 867	55. 576	41.006	34. 182	23. 206	45. 223
	HEMBA1001510	174. 254	120.414	343. 336	75.008	76. 932	73. 234	61.531	76.899
	HEMBA1001515	45. 158	26. 337	67. 169	15.756	15. 962	10.664	9. 567	12. 346
	HEMBA1001517	51,005	47.728	80. 287	34. 595	28. 246	21.020	17. 229	33. 972 4. 824
	HEMBA1001522	7.431	8. 980	7.032	7.566	5. 011 22. 395	6. 466 22. 767	6. 447 23. 530	17. 914
35	HEMBA1001526	48.774	21.300 85.570	32.732 262.800	18.831 70.163	46.649	44. 926	26. 457	37. 421
	HEMBA1001533	129. 423 59. 442	26. 656	27. 947	8.053	15. 558	53. 508	108.861	25. 371
	HEMBA 1001547 HEMBA 1001552	41.663	33. 242	115. 535	26. 222	30, 447	18. 258	21.358	25. 853
	HEMBA1001553	58, 388	75. 765	66. 228	32.264	36. 396	54. 513	64.874	41.905
	HEMBA1001557	182.516	80.827	161.852	69.344	80, 644	123.765	111.732	70. 946
	HEMBA1001563	39.649	31. 429	85. 246	26.057	12, 157	15. 987	10.085	17.083
40	HEMBA1001566	37.835	49. 954	108. 284	35.793	23. 255	25. 180	21.368	39. 375
	HEMBA1001569	75. 584	44. 631	109.624	35. 487	130, 340	63.130	44. 960	55. 257
	HEMBA1001570	198.300	125. 319	444. 153	119.332	74. 267	79.979	64.732	90.896
	HEMBA1001579	103, 128	60.654	48.704	22.469	22.629	67.058	24. 391	34. 300
	HEMBA1001581	153.698	126. 225		131.687	142. 104	91.884	67.267	94. 418
	HEMBA1001582	3, 551	7.087	15. 302	4.019	8. 190	4.888	4.671	5. 144
45	HEMBA1001585	27. 271	18. 375	25. 179	14. 108	5. 648	14. 993	7.628	12.297
	HEMBA1001589	109, 877	22.722	49. 216	20. 427	22. 904	64.665	57. 120	21.314
	HEMBA1001595	71.600	62.349	46. 938	34, 447	29. 362	34. 516	45. 233	35. 562 23. 101
	HEMBA1001604	41. 253	27.004	34. 167	15.004	6.061	21.932	18.414	15. 366
	HEMBA1001608	35.073	29. 270	41. 525	21. 276	22.867	22.699 81.914	42.373	58.652
50	HEMBA1001615	556. 575	105.703	103.519	47.685	62.056	54. 423	64. 359	36. 203
50	HEMBA1001620	134, 940	29. 972	79.824	31.924 15.048	19.545	42. 391	33. 266	40.516
	HEMBA1001621 HEMBA1001635	70.036	30. 704 29. 397	63. 807 35. 653	16.214	18.765	19.655	22.405	14.095
	HEMBA1001635	39. 932	18. 596	35. 798	14. 928	12.865	24. 352	31.819	22.414
	INCHORIUUIOJO	73.726	10. 330						
		40 403	45 102	1 79 599	1 78 457	1 77 444	1 75 101	311 11114	
	HEMBA1001640	48. 402	45. 105	79.588	35 084	25, 449	25. 101 48. 859	30.009 71.158	43.819
55		48, 402 82, 402 390, 307	45. 105 39. 456 66. 648	79. 588 75. 907 181. 929	28. 452 35. 084 51. 802	25. 220 112. 530	48.859 208.201	71.158	46. 463 96. 640

Table 19

HEMBA1001655	60.366	18. 983	58. 438	20, 404	25.072	27.162	29. 260	26.673
HEMBA1001658	6.754	15.270	17.542	13.420	5.060	4.800	4. 973	4.979
						24.098	22.916	24. 583
HEMBA1001661	87. 199	20. 304	32.793	13.066	8. 394			
HEMBA1001665	160. 583	20.830	54.460	12.363	48. 457	86. 024	73.847	21.248
HEMBA1001670	16. 953	38.651	17.002	34.999	14.855	17.849	22.906	29.478
HEMBA1001672	32.013	18. 885	29.000	10.798	7.763	13.782	17.314	12.393
HEMBA1001673	38. 188	67.401	34. 336	38.037	14.401	17.612	30. 520	43.461
HEMBA1001675	25. 652	15. 594	33.810	5.390	15.796	13, 173	20.020	12.830
HEMBA1001676	91.000	54.310	85.397	92.681	131.468	50. 365	47.230	68. 405
		128. 995	336.408	93.889	115.305	80. 843	48.879	83.933
HEMBA1001678	218. 382							
HEMBA1001680	82. 159	51.521	165.818	33.978	36.449	33. 368	38. 495	35. 261
HEMBA1001681	1.654	0. 785	0.840	2.142	2.581	2.772	2. 146	2.424
HEMBA1001684	143. 985	84, 151	377, 154	72.850	69.097	61.638	30.820	52.077
HEMBA1001695	16.068	10.112	14. 571	6.860	4. 930	4. 572	6. 164	7, 330
HEMBA1001702	26. 509	13. 637	8.186	8.466	4.041	2. 043	3.870	3.613
HEMBA1001709	67.279	26. 552	35.845	13.982	21.742	28.510	24. 540	19.603
HEMBA1001711	20.072	29. 559	39.037	20.902	21.639	12.713	14.718	33, 127
HEMBA1001712	80. 448	25. 222	51.628	19.393	12.482	38.014	39.474	14.831
HEMBA1001714	360. 368	55. 902	142, 225	33.748	51.048	144.094	124.654	59. 543
HEMBA1001717	78. 599	137.380	18. 549	12.298	5. 575	38.689	10. 120	6.047
HEMBA1001718	51.621	52.280	151. 597	31.305	21.166	29.146	14.075	24.411
HEMBA1001723	17.072	13.658	8. 525	5.653	8.811	9.350	11.097	7.268
		22.781	41, 531	15, 151	12. 421	15. 292	14. 020	16. 584
HEMBA1001731	35. 728							
HEMBA1001734	52. 546	40. 599	99.556	25.099	24.031	28. 537	17. 389	32. 936
HEMBA 100 1736	177. 269	58. 328_	110.046	33.820	58. 955	108.630	91.464	62.571
HEMBA1001741	41. 432	12, 649	29, 883	14.886	16.207	10.446	11.420	7. 286
HEMBA1001744	5. 531	6.849	12.961	13.191	14. 151	4.519	8.367	8.623
HEMBA1001745	41.752	17.786	36. 239	12.475	21.118	23.635	15. 410	16.514
HEMBA1001746	27. 437	14.874	24, 099	8.668	21.929	19. 488	11.306	10.070
HEMBA1001761	93. 148	46.911	179. 597	28.212	33.421	34.026	19.164	25. 901
HEMBA1001762	55.612	45.069	102.148	38.307	35. 260	33.316	21.274	45. 248
HEMBA1001781	13. 298	21.385	26, 693	6.898	17.098	52.601	11.768	23.068
HEMBA1001784	89. 965	43.765	70.064	26.575	31.708	50. 347	52. 265	31.618
HEMBA1001791	182.379	81.719	171.066	44.628	49.350	82.856	58. 215	48. 207
HEMBA1001794	248. 582	163.789	153,778	73.632	50.595	152, 279	178.827	132.329
HEMBA 1001800	23. 432	21, 165	27.668	11.281	20.728	24. 910	36.900	22.729
							6, 613	10.079
HEMBA1001803	17. 343	8. 333	22.801	6.620	6.043	7.560		
HEMBA1001804	109, 775	44.797	59. 456	29.337	34.849	44. 372	36.696	35. 851
HEMBA1001808	78, 129	23. 567	38.056	15.858	23.507	27.136	14. 673	12.332
HEMBA1001809	66. 887	31.733	54. 127	33.314	26. 179	35.618	41.552	46, 141
HEMBA1001811	58. 974	24. 196	37, 583	17.314	16.018	21.582	15.074	19.831
HEMBA1001815	71.286	63.775	155, 707	37. 153	29.944	35. 297	25. 257	24. 172
HEMBA1001816	38. 494	19.017	16.797	7.139	5. 598	16.061	22.304	14.646
HEMBA1001819	18. 590	21. 371	38.109	20.938	21.358	15. 313	14.917	25. 144
HEMBA1001820	10.884	9, 530	8.017	3.507	4,470	3.473	2.999	3.099
HEMBA1001822	74. 239	95.719	91.314	62.121	28. 285	42.988	38. 222	47.532
						89.045	64. 562	61.114
HEMBA1001824	155. 543	93, 583	301.248	95.135	67.478			
HEMBA1001835	23.616	7, 706	25. 753	5.777	19.660	19.809	12.020	10.462
HEMBA1001844	149.876	52.023	230.213	48.968	42.113	39.652	33. 559	40. 495
HEMBA1001847	52.045	19, 220	40,636	20.235	5. 196	35. 109	20. 186	35.814
HEMBA1001849	101.048	104.708	250. 547	53.025	28.022	40.644	33. 371	35. 250
							44. 306	
HEMBA1001850	105, 331	27.032	39.813	15.808	31.525	42.751		18. 213
HEMBA1001861	3. 104	4. 469	6.763	3.292	4. 454	2. 945	0. 995	3. 121
HEMBA1001862	50. 279	145. 708	102.412	25.750	34. 563	40.833	22.588	71.713
HEMBA1001864	24.313	31.572	50.378	32,237	24.991	21.182	21.031	28. 126
HEMBA1001866	57.711	54. 190	146.615	31.714	19.527	26.041	22.874	21.249
HEMBA1001869	55. 280	99, 559	58. 454	35.799		40. 562	22.644	40.891
HEMBA1001871	75.011	44. 336	77, 195	41.540	39.300	54. 584	34. 598	42.631
HEMBA1001876	34, 287	31.955	30.568	85.092	19.827	15.356	8. 554	21.861
HEMBA1001878	17. 361	17.619	17, 545		5. 481	11.657	14.965	18.117
HEMBA1001879	57.004	22. 429		16.562	20. 200	35.414	21.946	17.114
HEMBA1001884	68.009	84.640	41.930	38.470	27.460	36.604	25. 345	26.320
HEMBA1001886	12.711	12.605	37.824		15.893	14, 038	6.697	38.737
HEMBA1001888		46. 960	165. 623		21. 154	29.117	21. 131	33.090
ILEMOVIANI 1998	63. 251	1 40. 300	1 103.023	41.706	41.134	(3.117	1 (1.131	33.090

Table 20

	HEMBA1001890	42.902	42.848	42.779	30.112	25. 432	24. 430	22.605	26.730
	HEMBA1001896	66.448	24. 720	44. 103	21.972	17.708	30.703	19.628	23.571
5	HEMBA1001899	36. 251	25. 553	24. 121	14, 701	12.301	21.838	17. 455	20.813
5	HEMBA1001904	54.904	256.020	233.857	243. 646	55. 587	234. 548	188. 571	526.744
	HEMBA1001910	40. 309	10.865	13.738	11.244	8. 226	15. 357	15. 894	13.300
	HEMBA1001911	35. 962	23. 128	26. 357	25. 151	11.860	24. 224	22.870	18. 238
	HEMBA1001912	59. 924	66.966	97.679	51.180	45, 903	33. 336	33.019	40. 551
	HEMBA1001913	175. 368	39.664	67. 432	33.132	26. 376	63. 459	70.607	52. 824
10	HEMBA1001915	14. 756	14.666	30. 224	8. 295	7.629	17.718	6. 737	8. 522
	HEMBA1001918	5.018	8. 961	27. 591	7. 538	11.032	8. 265 7. 417	4. 852 7. 668	4.772 2.769
	HEMBA1001921	4. 431	8. 444	18. 196	11. 252	12. 587 3. 480	2. 935	1. 127	2. 898
	HEMBA1001931	3. 948	0.000	4.664 81,706	1. 422	16. 692	37. 223	29. 835	13.058
	HEMBA1001939 HEMBA1001940	94. 821 54. 512	24.679 33.931	145. 138	26. 273	27. 653	18. 649	13. 136	19, 514
	HEMBA1001942	38. 572	16.710	32.402	18.718	14. 782	25. 435	26. 410	16, 143
15	HEMBA1001944	210.898	71, 197	96.883	48, 156	38. 533	82. 132	92.097	74,740
	HEMBA1001945	31.531	17.019	14. 533	10. 175	3, 037	17. 421	12. 222	11.694
	HEMBA1001950	7. 103	7.424	9.611	3. 281	4. 091	7.632	5. 310	4,044
	HEMBA1001951	46.024	19, 234	101.026	19. 207	13. 212	23.714	20.006	19.402
	HEMBA1001958	44. 554	12.806	35. 277	17.321	13.181	22.652	28. 735	20.948
•	HEMBA1001960	20.513	7.802	16.888	8. 822	2.948	8.826	10.834	12.935
20	HEMBA1001962	4. 367	5. 104	4. 205	2.811	3.031	4.870	2. 364	2.994
	HEMBA1001964	35. 944	22. 281	62.761	18.757	6.661	17.775	· 8. 068	8.601
	HEMBA1001957	47. 345	29. 504	42.717	13. 525	22. 051	33. 555	23.601	37. 521
	HEMBA1001979	35, 138	6.478	16.732	12.797	5. 919	13.447	10. 355	9. 155
	HEMBA1001987	60.083	52. 275	190. 331 276. 566	45. 735 56. 455	24. 898 50, 862	25. 381 50. 789	17. 514 40. 252	28.891 54.919
25	HEMBA1001991 HEMBA1002003	111.286 66.389	79.833 23.989	53.710	17.039	17, 174	30. 547	28. 422	24, 474
	HEMBA1002005	86.885	41.457	150. 127	33. 935	15, 339	24. 541	24. 237	27, 345
	HEMBA1002008	32.101	25. 375	86.511	18. 349	8.912	7.593	18.519	14. 967
	HEMBA1002018	66.105	22. 380	36.174	16. 334	21.482	27. 922	34.098	27.804
	HEMBA1002022	13.986	8.018	13.490	0.000	2.985	5.730	6. 036	1.433
	HEMBA1002029	132.547	305. 823	115.974	144. 692	70. 087	74.071	37.046	204. 730
30	HEMBA1002030	17.077	10. 337	14. 524	5. 906	8. 466	5. 897	6. 258	6.824
	HEMBA1002035	48. 658	12. 959	10.324	14. 325	7.176	14.446	14.084	13.506
	HEMBA1002037	16. 343	34. 097	27.567 91.391	14. 451	12.568 8.370	15.087 6.020	13.819 17.500	12.140
	HEMBA1002038 HEMBA1002039	68. 477 15. 944	31.733	17.807	16.935	7, 910	3. 306	4.716	11.003
	HEMBA1002042	41. 657	27.877	32.654	21.111	14.815	10.217	24. 300	22.659
35	HEMBA1002042	149. 364	92.912	208.642	70. 906	53.861	84.089	81. 242	61.829
	HEMBA1002048	137. 253	29.889	60. 279	19. 894	21,605	66. 594	55. 483	30. 137
	HEMBA1002049	98.417	84.099	271.170	63. 157	87.434	48. 247	39. 557	53.676
	HEMBA1002053	33.636	19.194	25.821	11.890	16. 358	16.441	25. 376	27. 152
	HEMBA1002055	67.115	34. 916	39. 511	37.518	17. 449	25. 297	28.606	39.067
	HEMBA1002056	13.684	12.039	16.129	14.136	1.311	8. 564	4. 481	12.538
40	HEMBA1002061	11.815	14.960	29.478	10.168	10.973	11.179 39.904	9. 701 45. 653	8. 124 53. 581
	HEMBA 1002080	59. 350	80.319 7.502	81.497 15.981	7. 301	72. 416 10. 773	13.652	6.835	5. 555
	HEMBA1002084 HEMBA1002085	11.331 69.868	62, 174	111.196	13.760	19.083	101, 175	43, 117	14,011
	HEMBA1002092	127. 409	33.016	60. 924	24. 219	32.654	72.141	50.433	27,770
	HEMBA1002098	34.645	16.695	25. 357	15.741	15.632	18.082		20, 451
45	HENBA1002100	118, 301	90.733	129. 453	60. 276	41.079	89.713	44. 294	67.352
40	HEMBA1002101	57.160	69.427	106.418	34.067	32.565	38. 238	15. 932	74.139
	HEMBA1002102	104.746	76.058	178.766	45. 801	50.114	53. 399	40.628	54. 459
	HEMBA1002105	35. 380	25.812	31.300	14. 131	14.867	29.842	22.894	23.960
	HEMBA1002107	62.621	45.738	65.486	28. 199	31.808	52.057	163.850	77. 437
	HEMBA1002113	745. 018	396.517	1335. 986	321.385	369.500	391.825	236.013	348.025
50	HEMBA1002119	35. 812	23.546	72.351	18. 292	19.991	18. 086	26. 533	25.611 33.563
	HEMBA 1002125	84. 269	14.033 29.512	45. 440	15.858	29. 311	40. 381	28. 894 49. 691	37, 106
	HEMBA1002131 HEMBA1002133	37.736	19, 103	27.034	45. 990	11. 161	21.694	20.410	24, 305
	HEMBA1002133	25.756	10. 925		4. 978	11.839	9. 451	7.795	9, 431
	HEMBA1002141	20.036	14. 349		9. 608	3.638	14. 521	10. 225	10.190
55	HEMBA1002144	86.896	68. 335		56.749	45.612	36.918		

Table 21

	TIPUDA LABORA AT			22 200 1	40 410 1	46 210	67 057 1	02 212	45. 988
	HEMBA1002147	135.045	48.848	87. 208	42.412	46.318	67. 257	83.313	
	HEMBA1002150	347.113	89. 434	182. 502	48.715	86.270	215. 282	234. 394	85. 507
5	HEMBA1002151	60.410	19.140	11.868	10. 122	7. 938	26.996	19. 485	14. 196
ŭ	HEMBA1002153	32.258	25.478	35, 746	20. 325	25.638	15. 972	26.019	19.827
	HEMBA1002156	118.226	31.167	44. 382	21.446	21.743	47.426	40,620	16.858
	HEMBA1002160	166.654	114.853	336. 241	90.651	71.047	63.857	41.633	55. 419
	HEMBA1002161	72.851	68.019	132.156	42. 302	37.035	29.438	49.436	41.818
						51.141	55. 242	37.823	54. 952
	HEM8A1002162	122.516	52.989	307.464	68. 589				
10	HEMBA1002163	49.889	43.602	64. 932	20.426	7. 151	0.000	30. 979	32. 578
	HEMBA1002164	110. 221	59.044	71.408	32.652	19. 476	39. 440	43.639	52. 404
	HEMBA 1002166	312.686	256. 137	768.834	194.638	171.071	159.271	134.442	213.993
	HEMBA1002167	139.053	18. 430	69.693	11.057	30.091	50. 202	48.620	13.648
	HEMBA1002173	137.562	47.918	197.006	36.833	26. 164	29.478	20.169	23.078
	HEMBA1002177	100.895	25. 141	41.676	25. 857	17, 903	28. 153	22.687	14.081
	HEMBA1002178	102.831	19.500	46, 717	13.290	32. 323	37.856	44.115	27, 390
15	HEMBA1002179	55.617	56. 403	85.686	45.680	26.918	60.684	59, 110	64, 849
					43.915	27. 049	32.172	22.480	32.386
	HEMBA1002185	85. 236	71.958	212.844					18. 236
	HEMBA1002188	79.413	28. 280	31.826	23.275	21.094	33. 295	36.478	
	HEMBA1002189	56. 349	70.609	148.011	47.092	32, 460	30. 101	34. 751	30.532
	HEMBA 1002191	149.027	80.765	149.493	49.599	42. 372	60.095	35.614	44.348
20	HEMBA 1002 192	15. 125	24.996	24.821	15. 373	16. 495	12.778	5.075	13.566
20	HEMBA1002195	57.368	28. 535	52.828	16. 254	22.600	31.843	32. 995	29.791
	HEMBA1002196 .	14.884	12.040	36.633	16.632	15. 443	16. 808	12.691	17.451
	HEMBA1002199	24. 937	13, 539	27.878	15.728	17. 426	10. 539	19.664	8. 927
	HEMBA1002204	9, 525	5. 141	14.869	5.784	4.619	10.508	27.818	9,410
	HEMBA1002208	80.832	44, 154	68.317	68.994	37. 453	74.064	81.827	112.820
	HEMBA1002212	8.709	6.241	10.946	9.855	2, 602	5.864	5. 366	4.214
25	HEMBA1002215	36. 521	28.098	31. 165	19, 157	20. 170	17. 045	19.124	21.605
			62.759	64. 668	59. 460	28. 990	37. 379	29. 963	64.813
	HEMBA1002217	50.834				6. 409	5. 663	1.641	5.714
	HEMBA1002220	27.731	14. 997	21.655	8. 451				78. 924
	HEMBA1002226	91.222	113.507	269.906	85. 183	68. 283	59. 461	56. 996	
	HEMBA1002227	55.957	91. 527	79.169	45. 309	54.892	28.856	14. 142	101.597
	HEMBA1002229	170. 518	117. 589	418.739	112.916	121.703	85. 889	63.450	90.668
30	HEMBA1002237	47. 252	49. 329	124.721	32.838	24. 807	23. 399	15.399	26.185
	HEMBA1002239	103.363	107, 010	190.830	54.740	72. 381	50.451	45.873	70.581
	HEMBA1002241	70.729	45. 281	81.541	43.824	30. 449	54. 328	62.401	55.767
	HEMBA 1002253	25. 559	27.877	35.744	16.605	13.851	18.938	18.391	14.286
	HEMBA1002257	6. 344	5.787	15.404	4. 338	1. 225	7, 119	4.456	3.711
	HEN8A1002259	48. 436	19.578	38. 228	12.875	21, 884	23.928	18.619	17.988
35	HEMBA1002262	271.029	219.564	645. 284	192.491	147.403	112. 552	83.057	137.280
00	HEMBA1002265	56. 947	30.786	32.747	24.827	15.078	28.043	29.609	27.237
	HEMBA1002267	108. 413	102. 522	243.566	58.776	30.097	53, 750	24.099	29.752
						15. 579	28. 348	19.144	16.695
	HEMBA1002270	51.540	26.396	27.766	20.313		25. 959	14. 425	10.031
	HEMBA1002286	44.897	17.027	19.776	11.608	10.900			
	HEMBA1002290	46. 449	29. 289	34.095	19.879	8.778	26.461	22. 368	13.907
40	HEMBA1002302	152.883	48. 105	92.158	43.064	48. 204	66.899	80.872	58.027
	HEMBA1002304	6.050	6.814	19.492	7.905	4.038	7.098	5. 307	1.737
	HEMBA1002307	100. 402	132.737	29. 225	24.612	24.050	42. 355	39.076	37.573
	HEMBA1002316	504.772	93.620	191.534	46.814	134. 386	238. 599	265. 167	88.087
	HEMBA1002319	2.868	2.456	9.670	0. 933	4.715	4. 369	5.615	4. 579
	HEMBA1002320	10.783	7. 936	12.646	4.775	10.008	4. 330	5. 128	3.630
45	HEMBA1002321	10.743	9, 992	10. 165		2.547	7.952	4.048	5.700
45	HEMBA1002328	89. 382	28.578	41.753	17. 175	20. 280	46.772	34.722	18.301
	HEMBA1002333	63.542	21.208	32.148	11.559	15. 490	29. 410	33.449	21, 452
	HEMBA1002337		61.863	189.067	60.545	43.745	40.085	13.954	34. 456
		93.059		211.807	141.794	124. 733	173. 522	284.831	192.502
	HEMBA1002339	354. 195	154. 586				50. 321	45. 600	28. 278
	HEMBA1002341	116.488	29. 538	63.800		36. 228			
50	HEMBA1002348	6.882	4. 859			4.011	5.790	4. 476	4.606
	HEMBA1002349	6.318		13.603		2.590	6.088	1.306	3.748
	HEMBA1002353	14. 497	13,001	12.249		11.840	13. 977	17.141	16.760
	HEMBA1002356	104. 283	29. 278	40. 945	24.892	20.681	42.242	45. 108	28. 190
	HEMBA1002357	64.855		219.532		68.836	206.728	136.339	380.371
	HEMBA1002360	87. 281	64.882			56, 108	61.060	59.371	56.291
	HEMBA1002363	71, 449				43. 280	33.755	31.248	49.484
55	(IEMON100E303	1 . 1. 773	31.104	1 50.210	1 02.111	1 -3. 200			

Table 22

				- 2 52 1	2 175 1		4 146 1	10 000 1	0 205
	HEMBA1002365	13.435	10.346	9. 534	5. 175	9.470	4. 446	10.802	9. 325
	HEMBA1002370	29.997	4. 107	11.054	4. 163	3. 224	9.009	7.477	2.921
					15.862	10.204		29. 203	18.856
5	HEMBA1002374	91.498	18.475	11. 325			18. 275		
•	HEMBA1002376	186.416	75. 425	127.578	52.056	38.450	99. 590	90.190	86. 994
	HEMBA1002377	81, 350	41.908	63.893	37. 221	23.657	110.374	162, 166	50.770
•	HEMBA1002380	189.521	137.466	477.021	137.908	491.500	90. 431	81.778	127.767
	HEMBA1002381	195,037	101.891	447. 953	125, 938	88.330	90.756	70. 293	106.965
	HEMBA1002384	35. 247	22.319	42. 496	14.694	19.780	40.126	24. 243	12.399
	HEMBA1002389	44.796	8.467	36.790	11.793	9.362	18. 736	15. 497	20.728
10					16. 553	25.429	25. 964	22. 294	23.666
	HEMBA1002396	101.267	69.467	33.025					
	HEMBA1002402	75.818	24.148	28. 457	8.848	9.913	21.219	16.569	22.818
	HEMBA1002417	132.807	33.708	84, 436	22.910	38.826	58. 589	58.836	38. 486
	HEM8A1002419	75.547	31.202	41.590	13.558	16.457	27. 281	19.705	13.013
	HEMBA1002420	20.818	20.448	35. 559	17.034	13.878	23.652	14.721	24.637
							24, 230	61.011	21.849
45	HEMBA1002421	23.903	25. 285	59.023	7.957	14. 189			
15	HEMBA1002423	12.762	11.755	25, 941	12. 938	14, 177	14. 263	12.495	7. 512
	***************************************		32. 293	46.657	24. 424	25.667	42.797	41.513	31. 249
	HEMBA1002424	111, 995							
	HEMBA1002426	60.617	23. 489	45. 906	20. 305	25. 173	30.860	37.738	21.223
	HEMBA1002430	24.143	3. 128	4. 900	1.517	4. 594	3. 316	8. 552	3.069
	HEMBA1002439	59.808	37.476	93. 025	16.789_	23.324	47. 857	33.099	27.888
	HEMBA1002441	77.869	99, 262	110.341	38.723	34. 562	65. 309	85. 421	66. 581
20			15. 281	38. 384	7. 520	19.044	25. 972	22.845	22.015
	HEMBA1002454	58. 292							
	HEMBA1002458	57, 329	46. 103	101.242	30. 906	82. 184	61.800	25.094	59.039
	HEMBA1002460	32.814	9, 205	25. 085	12. 160	23.009	18.683	14.678	14. 249
								43.851	
	HEMBA1002462	98.420	38. 135	55. 208	10. 919	24. 257	49.697		32. 387
	HEMBA1002465	11.819	15. 260	28. 272	11.939	11. 225	10. 938	13. 593	20.635
		129, 538	61.348	120. 187	39. 999	39.213	76, 320	69.012	86. 309
	HEMBA1002469								
25	HEMBA1002475	3.180	5.116	4. 323	2.230	1.467	4. 495	10.058	15.691
	HEMBA1002477	93.696	64.730	238.114	55. 207	43.349	42.487	29. 532	52.786
								97.791	70, 487
	HEMBA1002480	210.023	58.823	84. 566	37. 478	45.060	106.554		
	HEMBA1002481	104. 499	76.474	222. 903	71.502	68.097	67. 421	42. 334	82.875
	HEMBA1002486	81.465	42, 269	169, 291	49. 953	40.852	39.475	29. 153	26. 233
									15, 278
	HEMBA1002490	66.695	11.331	31.314	14.602	25.852	35. 945	35. 954	
30	HEMBA1002495	59.387	12.315	25. 235	7.937	4. 091	17. 402	14. 269	10.773
			23.969	67.108	11.632	15.655	24. 420	8.272	12.219
	HEMBA1002498	56.425							
	HEMBA1002501	40.955	16.994	22.074	13. 575	16.498	21.707	39. 506	24.619
	HEMBA1002503	81.763	65.044	154. 595	39. 638	33.778	31.214	32.219	26.800
						120. 246	70.516	52.190	53. 323
	HEMBA1002504	155. 357	95.219	279.391	90.092				
	HEMBA1002508	99.443	88. 234	259.961	107.085	79.039	59. 181	59.924	61.423
	HEMBA1002513	50.560	22. 902	30.431	26. 184	20, 783	30, 500	32, 903	22.864
35									
	HEMBA1002515	60. 938	23.064	25, 098	16. 172	5.716	20. 264	20.643	13.727
	HEMBA1002524	94. 350	36.789	56.675	25. 998	28.978	49.840	57. 148	25. 205
			19.632	26.764	12.798	20. 203	16. 422	17. 588	15.759
	HEMBA1002538	116.609							
	HEMBA1002542	81.641	81.952	188.888	54. 986	41.864	32.890	30.719	38. 321
	HEMBA1002544	52.394	49.175	98. 415	47. 569	28.375	28.766	20.948	21.614
		76. 538	62.763	156.051	47.625	74.374	45. 975	34.756	46.753
40	HEMBA1002546								
	HEMBA1002547	11.448	4. 516	10.647	4.733	12,220	11.801	9, 959	7. 127
	HEMBA 1002550	67.373	39. 322	48. 468	15.671	16.497	121.814	94. 586	25. 401
				27.085	11.976	8.787	41.811	16.656	18, 665
	HEMBA1002551	94. 391	14. 109						
	HEMBA1002552	204. 583	77. 430	205. 444	49. 448	44.756	67.408	63.216	57.684
	HEMBA1002555	25, 583	16.987	6.743	7.020	5.608	14, 795	10,111	7.416
45	HENBA1002558	92.744	77. 405	245. 703	59.079	41.247	33. 253	41.517	41.270
73	HEMBA1002561	53, 810	51.725	155.895	34. 956	27.689	17. 264	10.138	27.124
				15. 435	8. 259	18.723	12.036	9, 056	10. 429
	HENBA 1 002562	15, 261	10.822						
	HEMBA1002568	24. 946	17.442	35. 354	17. 552	10.576	15. 262	16. 158	22. 328
	HEMBA1002569	112.340	34, 133	118.192	37.823	57.431	54. 936	26. 164	27, 309
	HEMBA1002570	43. 528	50.809	52.195	34. 901	23.728	28.874	9.812	50. 494
	HEMBA1002574	106, 101	25, 148	46. 793	16.369	26. 322	57. 278	42. 795	31.310
50	HEMBA1002583		17.582			13.418	20. 158	16.837	16.418
		35.042							
	HEMBA1002587	61,527	32. 123	45.811	22. 217	18. 974	32.461	40. 250	39. 915
	HEMBA1002590	151.583	106.074	287.276	84.766	32. 321	58. 221	38.642	53.855
	HEMBA1002592	97. 854	85. 949	220. 496		52.684	53. 553	35, 724	57. 578
	HEMBA1002595	146,016	25. 688	60, 427	24. 156	31, 909	86.770	79.174	26.760
			41.926	56, 054		35.650	35. 839		47.074
55	HEMBA1002609	97. 442	1 41. 370	30, 034	1 69.461	1 33.030	1 33.039	1 44.008	1 41.014
5 5									

Table 23

		00 200 1	44 417 1	FO. 446 I	72 077	12 000 1	21 055 1	10 012 1	40 126
	HEMBA1002617	26. 792	86.617	59. 446	73. 277	12.909	21.055	16.612	49.136
	HEMBA1002619	101, 131	25.998	30. 959	15.921	21.913	40.814	35.003	28. 108
5	HEMBA1002621	14. 592	25.845	18. 082	8. 927	7.391	8.869	5.823	12. 283
3	HEMBA 1002624	254.635	42.837	73.568	48.036	71.673	113. 228	101.786	53.514
	HEMBA1002528	13.044	21.509	23. 549	9.956	16.559	10. 257	7.527	11.624
	HEMBA 1002629	32, 199	16.370	29. 306	15.884	5.722	15.410	42.964	19.680
	HEMBA1002632	55. 206	48.044	90. 986	36.904	27.840	28.811	37.912	40.048
	HEMBA1002645	95. 909	89.897	220. 184	68.171	48.643	56.847	41.355	59. 567
	HEMBA1002651	39.882	27.730	33.313	16.958	11.617	23.904	29.214	16.599
10	HEMBA1002652	107.869	24. 187	46.646	22.248	22, 950	37.216	25.827	23. 282
	HEMBA1002659	133. 320	62.916	259.854	57.860	53. 172	46.511	45. 193	47.291
	HEMBA 1002661	88.495	68.014	154. 170	35. 196	22. 499	26.290	22.314	23.727
	HEMBA1002666	34.174	20.511	39.391	17.036	15.852	20.842	19. 202	13. 470
	HEMBA1002667	155. 384	166. 244	164.658	29.523	520.013	30.234	25.612	83.769
	HEMBA1002673	71.650	40.718	73.822	33.403	39.914	40.129	38.619	22.532
15	HEMBA1002678	161.681	89. 986	247.534	84.722	54. 176	46.941	51.944	77.085
	HEMBA1002679	56.416	61.838	66.537	37.679	18. 172	29.420	38.238	44.113
	HEMBA 1002688	6, 756	3.364	5. 387	3.816	1.793	4.608	3.600	2.944
	HEMBA1002696	49.639	17.555,	29, 241	14.788	12.463	31.752	34, 100	14.772
	HEMBA1002703	185, 328	96.718	97, 793	54.473	50.688	113.980	87.727	59.878
	HEMBA1002706	49. 533	30, 340	35. 679	18.469	19.118	26.777	29.277	29. 224
20	HEMBA1002712	52.878	59, 111	110.506	41.591	43.597	39.604	30.872	26.457
	HEMBA1002715	149.045	59.858	87.643	47.473	41.264	95.279	127, 808	65.580
	HEMBA1002716	23.142	6. 155	17.077	15.783	23. 557	19.064	27.647	7.572
	HEMBA1002718	26. 328	19.063	41,749	26.345	16.735	28. 367	26.822	21.779
	HEMBA1002728	117.984	88.950	293.019	81.290	43.679	65. 830	46. 321	57.003
	HEMBA1002730	131.726	26.862	67.877	28.628	36.686	49. 987	50, 380	43, 208
25	HEMBA1002734	77.679	26.481	34.604	21.128	21.756	41.413	60.057	45.992
	HEMBA1002742	10.730	11.276	12.768	7.910	1.394	8. 502	8.297	10.909
	HEMBA1002746	60.876	22.803	35, 400	15.830	15.630	30.605	31.889	32.759
	HEMBA 1002748	76,748	26.130	38.569	17.760	32, 833	43, 493	53, 440	49.691
	HEMBA 1002750	40.663	45. 306	95, 205	18, 200	10.037	22, 527	29.331	30,774
	HEMBA 1002755	94.758	62.505	220.964	63.414	37. 572	44, 593	28. 497	39.737
30	HEMBA 1002759	13. 935	3, 117	8. 450	3.792	2.291	8.714	10.261	5. 285
30	HEMBA1002763	430. 941	88. 931	172.920	71.623	88. 921	195. 471	197, 995	118, 224
	HEMBA1002767	65.682	25. 272	35.782	14.035	19.183	31.497	33. 393	18.347
	HEMBA 1002768	100, 803	57.554	59. 457	35.570	28.006	43,770	40.930	38.215
	HEMBA 1002769	103. 210	30. 236	54.098	17.099	19.753	35.636	41.922	26, 940
	HEMBA 1002770	20, 350	16.268	28.054	21.736	10.754	12.030	14.991	11.776
ne.	HEMBA 1002777	130.615	37.655	72.072	41.794	31, 219	54.881	59.342	43.652
35	HEMBA1002779	97.457	29. 259	75.705	22.719	22,643	33, 689	38. 357	27.804
	HEMBA 1002780	72.338	50.411	181.356	42.070	19.957	31.370	27.642	39.672
	HEMBA1002790	87, 371	61.291	152.514	38.033	29.616	28.032	20.352	34.761
	HEMBA1002794	202.405	77.515	95. 182	31.252	41.834	100.167	80. 301	50.036
	HEMBA1002798	9, 194	21.334	22.468	20.281	12.823	11.156	11.647	15.735
	HEMBA 1002801	10.311	4.603	11.704	3.190	4. 420	3.016	13.829	6.693
40	HEMBA1002810	42.583	45.313	55.088	35.416	29.480	60.935	44.046	51.794
	HEMBA1002816	52.084	37.823	56.994	35.902	25.574	33.389	50.974	49.045
	HEMBA 1002818	321.516	100.826	187.799	84.893	81.695	152, 339	171.186	117.409
	HEMBA1002820	139, 924	107, 278	533.137	90.533	79.745	59.869	54. 302	52.958
	HEMBA 1002826	40.776	6.495	16.825	5.349	3.319	11.765	7.355	8.363
	HEMBA 1002833	119. 102	44.248	40.839	17.864	23.748	44. 398	57.302	36.668
45	HEMBA 1002850	5. 941	8, 407	13. 251	6.179	2.932	4. 352	4.844	3.735
	HEMBA1002862	60.735	32.524	30.030	9. 593	9. 527	27. 595	19.397	18, 101
	HEMBA1002863	77. 126	30.401	44.872	22.577	28. 639	50. 264	55.374	45.005
	HEMBA1002867	25. 385	13.583	42.122	15.283	9. 501	22.992	15. 180	16.196
	HEMBA1002876	101.249	55.603	38.073	36.480	23.017	53.318	51.363	56.689
	HEMBA1002886	9.474	14.188	23.688	7.657	11.980	14.640	6.432	18.574
50	HEMBA1002896	78.580	27. 420	49.774	16.754	20.365	36.684	35.283	42.662
	HEMBA1002913	126.001	32.845	58.138	14.590	22.846	54.873	56.608	38.801
	HEMBA1002921	63.378	25.443	37.615	15. 333	19.054	28.881	37.595	34. 298
	HEMBA1002924	65, 007	29.109	104. 125	15.411	19. 920	31.099	23.998	19. 182
	HEMBA1002934	432.841	308. 291	644. 522	180. 470	145. 293	273.733	166.153	242.809
	HEMBA1002935	92.005	52. 184	221.722	49.477	41.867	34. 331	29.646	38.889
55	LICENVIONE 322	1 32.003	1 45. 107	1 451.166	1 70.711	1 001	1 -7. 04.	1	
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Table 24

	HEMBA1002937	38.698	30, 844	33.817	12.784	18. 251	14, 107	24. 131	18.662
	HEMBA1002939	39.755	22.867	33.838	19.077	13, 734	19. 266	17.364	17.750
	HEMBA1002944	53, 762	33. 349	51.861	21.860	18. 241	23.920	21.112	16.286
5	HEMBA1002944	38.716	29. 783	39, 196	19.808	29.614	19.702	28. 422	21.177
	HEMBA1002954	24. 907	8. 542	20.941	9. 265	13.758	15.056	7. 297	13. 424
	HEMBA1002962	86, 680	62.578	220. 246	62.027	37.753	44. 037	31.812	41.725
			78. 850	221.414	65. 545	40. 380	43.093	38.816	50. 281
	HEMBA1002968 HEMBA1002970	105. 871 48. 034	34.741	30. 834	18.482	6, 639	17. 125	23.514	36. 180
				35.618	25.614	12. 932	25. 193	14.823	23. 202
10	HEMBA1002971	39. 492	44. 145 70. 965	156. 167	43.307	28. 902	29.947	26, 101	34. 769
	HEMBA1002973	83.710	19. 362	27.056	13.075	20. 398	11.324	16.059	13. 956
	HEMBA1002978	35. 833	35, 200	56.576	23.695	26, 105	33. 054	37. 199	21.249
	HEMBA1002981	107. 112		116.532	27. 950	26, 158	37.462	28. 927	20. 335
	HEMBA1002985	79. 217	44. 154		49.967	64. 529	38. 333	28. 919	20. 529
	HEMBA1002986	61.056	78. 203	68.834 71.802	20.621	8. 965	16. 229	15. 956	22.796
15	HEMBA 1002 988	37. 307	36.609 72.656	79.841	50. 454	34. 289	57.004	61.291	91, 211
	HEMBA1002992	97.720		55. 081	36. 903	25.007	38.630	19.510	48. 529
	HEMBA1002995	51.473 41.734	63.779 70.805	29. 264	27.019	33.664	24. 201	18. 442	25. 973
	HEMBA1002997			18. 357	11.146	7.034	12.086	13. 966	9. 970
	HEMBA1002999	35. 341	16. 456	35, 194	15.119	16. 204	20. 866	27. 891	20.055
	HEMBA1003004 HEMBA1003006	55, 654 40, 682	33.689 24.886	20,750	20.903	26. 595	25. 445	20.310	20. 924
20	HEMBA1003008	29. 269	20. 922	74.697	25.061	17.787	10. 271	5. 688	12.638
	HEMBA1003021	130.889	123.646	311. 225	101.957	95. 443	64.844	60, 969	90. 296
	HEMBA1003027	54. 935	32.610	44.710	18.890	52. 131	26. 286	28.112	31.561
	HEMBA1003029	33. 333	42.436	60. 787	20.829	34.111	29.704	49.230	45. 833
	HEMBA1003031	34.000	25. 311	18. 494	14.998	13, 316	13.955	15.773	27.136
	HEMBA1003032	171.114	46.990	71.365	23.640	50.526	81.278	84.036	46. 352
25	HEMBA1003033	168. 563	118.674	378.771	109. 222	90.670	70.150	55.336	77.819
	HEMBA1003034	173.162	127.221	484.135	108. 238	85. 630	61.733	36.799	63. 312
	HEMBA1003035	11.693	5. 195	9.305	4.478	5. 058	11.024	2.553	4. 409
	HEMBA1003037	261.159	89.481	145.321	58. 521	65.732	104.677	89. 571	71.674
	HEMBA1003041	103.945	105.085	291.931	93.188	75. 193	53. 097	39. 564	58. 217
	HEMBA1003046	40. 254	39. 965	46.856	26.192	11.615	35.659	25. 378	32.416
30	HEMBA1003047	127.888	49. 341	139.750	32.219	32. 320	57.450	33. 390	28.702
	HEMBA1003048	87. 433	35. 962	42.305	12.040	20.442	39, 108	29. 597	21.461
	HEMBA1003064	6.366	8.535	6.201	8.809	4.415	7. 239	3.330	7.829
	HEMBA1003067	55.833	34. 508	77.097	26. 154	20.523	28.755	24.783	17.488
	HEMBA1003071	54.728	22.509	28, 869	17. 461	19.647	20.624	22.285	19.438
	HEMBA1003072	62.421	30.769	31.225	26. 146	22. 906	21.483	17.616	19. 134
35	HEMBA1003076	111.254	51.085	78. 972	37. 151	40.422	49.911	47.023	64. 737
	HEMBA1003077	36.471	15.407	24. 522	8.009	8. 453	18.661	13.797	5. 837
	HEMBA1003078	34. 143	38.741	77.906	31.907	37. 169	17.933	17. 439	18. 923
	HEMBA1003079	28. 559	39. 563	41.646	26.110	25. 889	25.576	18.026 29.619	24. 526 67. 469
	HEMBA1003083	61.036	48. 635	169.439	52.788	60.016 12.063	41.611		
	HEMBA1003086	49.032	40.488	154. 409	29.869		15.848	16.039 25.027	19.219
40	HEMBA1003090 HEMBA1003094	34.778 184.999	14.860 43.363	23.758 72.116	12.710 30.096	24. 132 53. 636	78.251	84. 551	34.775
		31, 440	18.030	25.774	10.290	11.781	14.033	27.791	11.348
	HEMBA1003096 HEMBA1003098	36, 774	64. 970	88. 562	34.074	24. 271	25.656	18.003	31.059
	HEMBA1003101	55.716	24, 121	22.316	11.682	13. 163	21.315	25.117	15.689
	HEMBA1003109	48.411	21.093	39. 285	21.315	21.724	27.826	31.034	21.809
	HEMBA1003114	41, 101	24.786	22.792	14. 164	14.657	18.320	15. 152	16. 038
45	HEMBA1003117	22. 939	13. 535	20, 191	6.812	10. 538	14.917	18.015	12.566
	HEMBA1003120	24. 531	24.408	55, 805	26. 574	13.838	15.423	15.080	21.728
	HENBA1003129	40. 276	46. 792	104, 463	37.995	37.989	21.990	26.267	38. 207
	HEMBA1003133	50.080	22.873	35.022	15.164	20,000	21.592	25. 551	27.656
	HEMBA1003136	146, 630	23.706	65, 990	18. 301	7 31.049	69.754	51.669	25. 346
	HEMBA1003142	69.008	47.867	130.557	32.955	30. 384	25. 274	27.118	29. 493
50	HEMBA 1003 148	59. 282	20.084	32,740	18. 292	18. 973	32.206	22.003	24.674
	HEMBA1003151	53.856	20.003	51.824	13. 233	9. 854	27.114	22. 251	13.546
	HEMBA1003152	20.577	9.803	19. 388	10.017	5.761	31.586	23.227	6.853
	HEMBA1003157	16.477	9, 272	16. 246	9,919	17.605	7.547	10.156	10. 181
	HEMBA1003168	293.814	257. 380		260. 521	221. 325	137. 459	148.208	199.758
	HEMBA1003171	17. 730	8.702		6.499	6.963	7.361	5.733	7.164
55	[HE-041003111	1 11.190		1	1 -1 100	, 3,000	1		

Table 25

								13-106-	15 625
	HEMBA1003175	38.620	40. 445	100.302	29.594	17.624	21.152	13.386	15. 936
	HEMBA1003179	63. 835	33.869	50.631	27. 163	25. 502	35.500	39.052	37.713
5	HEMBA1003186	100.461	75.611	231.787	75.781	58.278	54. 222	55. 862	61.615
•	HEMBA1003196	36. 422	27, 557	45, 633	20.623	18.740	21.756	30. 501	35.864
	HEMBA1003197	8. 462	9, 564	5. 534	5. 965	4.051	3.138	7.054	7.066
				81. 183	15.696	16.799	9. 492	17. 381	15. 917
	HEMBA1003199	34.650	18. 409						
	HEMBA1003202	79. 337	59.764	236.822	43, 286	41.820	31.106	32.936	45. 183
	HEMBA1003204	66. 523	56.272	172.818	48.560	31, 451	33.193	27.421	28. 849
10	HEMBA1003210	23.713	52.768	35. 498	5. 529	38. 451	16.353	59. 417	17. 563
,,	HEMBA1003212	126. 394	90.709	372.474	74.164	62.392	59.663	45.714	54. 363
	HEMBA1003218	19.415	13, 105	13.670	6.371	4.792	13.681	10.789	6. 536
	HEMBA1003220	81. 171	86.642	147. 453	89. 495	42.391	47. 586	54.647	123.019
		25. 803	22. 891	28. 577	7. 994	11. 404	10.413	9.856	14. 985
	HEMBA1003222					20. 280	48. 243	44. 574	19. 547
	HEMBA1003225	105. 735	21.238	40.848	11.586				
15	HEMBA1003229	30.394	26. 363	41.333	22.998	17. 475	14.707	20. 154	19.749
	HEMBA1003230	69. 643	70.015	42.439	31, 176	20.775	56.815	40. 191	75. 238
	HEMBA1003235	44. 989	43. 337	105. 267	33.038	19, 405	20.834	22.018	29.856
	HEMBA1003236	8.677	17.896	8, 735	7.270	7. 328	17. 286	5. 295	18.441
	HEMBA1003250	7. 260	12.5981	12.993	4.750	4.815	7.242	5. 982	4. 378
	HEMBA1003252	56.274	51.495	65. 197	28. 241	33.512	44.917	62.506	60.076
	HEMBA1003257	71.751	16. 083	40.414	13. 391	19.441	38. 988	28.614	19.028
20				46, 948	14. 167	12.769	11. 524	8. 622	17. 414
	HEMBA1003268	19. 492	18. 996	40. 346					
	HEMBA1003273	48. 113	38. 933	125. 242	29.404	21.135	22.989	17. 240	24.704
	HEMBA1003276	36. 279	34.802	113, 584	23.812	17. 208	20. 437	14. 685	26. 145
	HEM8A1003277	31.363	12.827	21.514	10. 462	11.287	13.206	16. 182	14. 465
	HEMBA1003278	36.998	24. 906	71. 222	17.479	15.791	16. 787	10. 948	17, 841
	HEMBA1003280	50.716	16,000	38. 057	15.933	20.792	37. 901	30.931	31. 493
25	HEMBA1003281	66.732	21. 393	32.728	15.032	18.415	26.844	28. 577	24. 898
	HEMBA1003284	9.746	8. 482	12.941	5.779	5.747	5.813	3. 545	3.499
	HEMBA1003286	69. 502	35. 947	60.729	21.827	29, 473	52.233	50. 283	47.695
	HEMBA1003291	13. 248	9. 951	10.909	3.504	18, 100	6.561	6.341	7, 647
	HEMBA1003294	69. 599	52, 239	168.555	39. 127	38.460	40.377	24.057	27. 486
	HEMBA1003296	61. 933	31.456	37.947	21.206	23. 199	23. 249	34.580	37, 768
30			5. 972	8. 976	6. 154	8.839	4. 199	3. 461	3. 227
30	HEMBA1003304	7.117				11. 282	8. 448	6.333	9. 387
	HEMBA1003306	17. 590	15. 590	22.443	8.410				
	HEMBA1003309	6.845	10. 103	12. 198	14.015	7.776	8.709	3.955	18. 326
	HEMBA1003314	637.052	210.608	238.618	105.098	198. 106	299.884	273.738	171.516
	HEM8A1003315	83. 736	51.612	84. 690	32.381	29. 482	56.694	53. 105	54. 024
	HENBA1003322	108. 401	88. 539	256. 570	51.502	51.083	44. 130_	42.804	45. 519
35	HEMBA1003326	42.723	20. 581	14.759	11.799	7.780	18.087	12.420	9. 516
	HEMBA1003327	61.811	36.702	87.698	28. 181	19.784	18. 596	17. 453	18. 377
	HEMBA1003328	53. 406	51.712	114.941	36.926	25.000	18.669	22.079	32.865
	HEMBA1003330	108. 955	82.099	207.708	73.413	52.244	50.838	55.920	55. 390
	HEMBA1003348	121. 625	110. 275	337. 182	94, 209	99,717	67.000	43.513	80.023
	HEMBA1003369	5.861	23.644	14. 930	4.979	1.726	9.064	3.020	5. 373
		315.016	197.956	369.117	140.044	139. 216	140.758	150. 458	124. 948
40	HEMBA1003370 HEMBA1003373		31.291	53. 330	17.430	5.513	19, 164	8.117	19. 638
		50.135		519.668	126.099	89.798	108. 226	81.818	107. 084
	HEMBA1003376	174. 269	170.290						
	HEMBA1003380	43.015	24.657	74.071	29.281	24. 407	19.711	13.485	20. 047
	HEMBA1003384	25. 555	30.071	68.079	15. 389	9.455	11.810	8.800	14. 281
	HEMBA1003387	6.515	2, 588	2. 697	1.577	1.109	1.803	1.986	3. 464
45	HEMBA1003392	111. 457	25.882	42. 253	17.323	29.007	50.086	29.337	23, 550
45	HEMBA 1003395	16.068	18.666	35.483	15. 254	9.873	10. 355	6. 207	12.514
	HEMBA1003399	45. 227	21.480	37.035	19. 231	15.354	19.471	27.860	34. 116
	HEMBA1003400	116, 210	36, 907	58.706	24.811	49, 133	53.819	60.041	53, 109
	HEMBA1003402	32.500	16. 239	27.864	8.795	12.867	17.141	11.617	14. 596
				46.720	20. 221	26, 579	36.738	44. 891	45, 870
	HEMBA1003403	60. 260	43. 377				84. 358		
50	HEMBA 1003408	196.676	49.687	70.460	29.354	50. 910		77.062	46. 433
	HEMBA 1003412	104, 813	43. 934	55.699	47. 250	43.763	61.953	59.463	47. 139
	HEMBA1003417	22. 445	13.970	25.036	8. 433	7. 282	10.593	5. 696	11.032
	HEMBA1003418	57.411	57.397	76. 232	97.795	45. 336	43, 450	22. 206	90.604
	HEMBA 1003420	29.838	15.856	201.831	11.319	8.067	11.379	12.938	14. 721
			15, 895	21.652		6,723	8. 483	10.838	9. 083
	HEMBA 1003425	1 17.400	1 13.033	1 61.002	4.133	0.123	_ 0.463	10.030	J. 003
	HEMBA 1003425	17. 466					10.074	11.092	11. 581
55	HEMBA1003425 HEMBA1003433	23. 931	18. 435	24. 576	12. 136	10. 421			

Table 26

					10 300	00 041	45 000 T	31 630 1	15 045
	HEMBA1003440	91.727	41.727	39. 257	19.755	26. 941	45. 998	31.620	35.845
	HEMBA1003442	7. 090	22. 535	10. 452	33.897	10. 259	15.118	7.093	14.790
5	HEMBA1003447	82.161	36.670	48. 248	26.789	18. 587	41. 591	42.314	35.065
•	HEMBA1003453	50. 472	26.692	25. 954	16. 130	11. 252	16.584	28. 534	21.256
	HEMBA1003461	55. 687	25. 328	42.686	17. 261	18.856	27. 281	22.795	17.854
	HEMBA1003463	40.102	23.311	34. 469	13. 456	19.704	20. 277	16.984	18. 124
	HEMBA 1003465	92. 245	40. 963	61.816	28. 410	36.051	39, 389	40. 220	36.851
	HEMBA1003480	114.075	114.841	266.076	76. 366	67.942	56. 459	51. 589	62.191
40	HEMBA1003485	44. 403	28.836	33.659	14. 371	8. 636	26. 284	16.036	14. 582
10	HEMBA1003487	42. 939	15.463	23.730	9.752	15.729	24. 902	21. 136	16.494
	HEMBA 1003492	31.026	21. 538	56.674	14. 934	12.014	12.082	9. 567	14.655
	HEMBA1003494	97. 366	260. 496	50.174	48. 821	12,504	74. 554	20.623	180.841
	HEMBA1003497	39.000	17.943	24.659	11. 432	13.881	21.376	18. 562	6.072
	HEMBA1003503	54.774	21.485	28. 175	12.948	17.154	30.911	36.463	16.806
	HEMBA1003511	18.672	14.740	43.023	11.794	13.330	8. 925	16.405	11.615
15	HEMBA1003528	385. 123	191.234	239.319	81.329	123.915	213.945	179.430	96.672
	HEMBA 1003530	43.820	12.384	23.693	10.695	21.216	20.067	28.030	16. 204
	HEMBA1003531	111, 104	73.542	215.578	67.833	214.022	56. 139	50.217	66. 992_
	HEMBA1003532	145, 137	62. 379,	83.827	37. 506	53.388	90.314	77.728	60.515
	HEMBA1003538	61.123	20.746	32.949	11.160	19. 286	34. 305	28. 231	13.837
	HEMBA1003545	21.489	10.501	20.608	5. 904	7. 197	10.239	6.617	8.168
20	HEMBA1003546	31.371	32. 365	28.613	13. 365	226. 243	16.427	16.554	24.821
	HEMBA1003548	4. 466	8. 124	9.845	4. 563	7.542	6.155	5.647	8:387
	HEMBA 1003553	79.837	51.515	50.379	23. 327	28. 564	49. 154	63. 525	48. 955
	HEMBA1003555	20.066	8.873	13.692	4.762	3. 584	10.112	10.962	6. 521
	HENBA1003556	57. 280	36.399	128.391	29. 283	16.426	19.257	18. 121	24.622
	HEMBA1003560	9. 290	4.426	2.529	2.848	1.767	2.983	6. 207	6. 539
25	HEMBA1003565	42, 648	29.588	20.996	8. 344	13. 984	21.927	21.847	22.043
	HEMBA1003558	7.244	1.649	7.712	2. 430	3.763	3. 172	2.836	2. 592
	HEMBA1003569	25. 048	20.536	23.764	33. 957	13.740	16.235	19.512	16.518
	HEMBA1003571	111.721	94. 378	326.335	84. 368	71.788	50.029	48.011	59, 960
	HEMBA1003579	3. 335	7. 399	15. 353	6. 553	8. 948	2.872	9. 198	6. 421
	HENBA1003580	274, 105	50. 292	102.103	26.686	59.875	128.943	110.375	35. 695
30	HEMBA1003581	112.013	31.295	94.083	21.641	36. 215	54.336	50.711	21.238
	HEMBA1003591	97.076	64.326	77. 160	89.876	47.882	53.615	40.656	45. 172
	HENBA1003595	32.697	22.842	84.629	19.075	11.339	6.305	5. 581	18. 085
	HEMBA1003597	48. 561	25.846	108.491	20.931	15.952	19.375	17.580	20, 153
	HEMBA1003598	49.728	20.134	22.468	12.142	11.688	18.934	21.743	15.025
	HEMBA 1 00 3 6 0 0	32.772	35.099	56.905	26.268	29. 290	38.873	53. 305	56. 783
35	HEMBA1003602	18. 248	10.116	16.162	6. 182	10.970	8.064	14.736	17. 188
••	HEMBA1003604	205. 949	53. 579	69.723	24. 549	49. 902	105.181	98. 166	47. 144
	HEMBA1003610	140.996	29. 255	95.048	15, 492	103.150	72.233	54.670	30.688
	HEMBA1003615	57. 258	20.035	34. 102	12.808	16.022	24.378	18, 759	20.876
	HEMBA 1003617	48.414	20. 375	29, 789	12.148	22. 291	18. 199	18.770	18. 242
	HEMBA1003620	52.899	22.318	45. 502	19.575	19.952	25. 239	39.072	29. 451
40	HEMBA1003621	102.827	102.094	226. 373	80.194	64.742	58.874	67.142	60.680 13.833
.•	HEMBA 1003622	19.815	13.838	25.009	16.055	8. 339	12. 261	15. 369	
	HEMBA 1003630	20.008	16.381	30. 244 106. 379	23. 251	5. 573 18. 468	9.992	10. 303	11.422
	HEMBA1003637	37.880	29.848						20. 420
	HENBA 1003640	39.068	31.672	100, 901	22. 572	22. 223	12,142	17. 417 54. 230	5.711
	HEMBA1003645	25.820	19, 380	48. 445	13.481	9. 247		19, 177	19. 938
45	HEMBA 1003646	38.243	16.329	22.003	9.624	7. 892	7,607	7. 882	10.058
70	HEMBA1003647	10.261	10,718	12.323	7.860 28.981	19. 429	18.898	23, 172	30.178
	HEMBA1003656	40.171	31.269	66.874 19.352	6. 387	10.041	10.909	14.055	18.544
	HEMBA1003662	25. 325	17.011		5. 803	8. 262	9.774	15. 332	13.851
	HEMBA 1003666	23.086	11.187	17.407		131.792	179.317	140.769	174. 256
	HEMBA 1003667	304. 975	209. 929		96.636	2. 565	7.057	6. 425	7.073
50	HEMBA1003670	12.944	8.894	15. 235	3.344	62.734	66.675	65. 424	47. 173
50	HEMBA1003674	143. 262	32.196 45.946	51.919	33.863 45.985	43.474	38.916	30. 594	46.808
	HEMBA1003677	80.516		220.695		5. 941	12.433	12.034	11.720
	HEMBA1003679	25. 325	7.795		6.727	23. 985	25, 419	38. 990	39.343
	HEMBA1003680	42.317	25.723 10.175					10. 182	10.574
	HEMBA1003684	18.273	65. 531				71.797	85. 431	56.592
	HEMBA1003690	115.021	03.331	1 13.070	1 40. 324	1 43.039	1 11.131	1 00.401	1 00, 332
55									

Table 27

	HEMBA1003692	83. 253	96. 347	194. 372	63. 188	40.872	47. 354	39. 288	42.644
						29.023		46.956	36.550
	HEMBA1003702	88. 125	35. 028	48. 251	23.719		42.879		
_	HEMBA1003711	93. 732	50, 280	140, 199	32.886	33.424	47, 500	41.959	36.807
5						28. 237			
	HEMBA1003714	75. 923	20.696	37. 340	14.414	20. 231	32. 029	29. 145	16. 214
	HEM8A1003715	54. 160	54. 486	142.871	31.894	31. 122	28. 832	20.640	26.672
	HEMBA1003717	70. 553	38. 574	120. 922	45. 101	29. 491	29. 344	27. 200	38.418
	HEMBA1003720	83. 687	94. 829	133. 285	55, 896	49.519	43. 330	22, 099	41. 137
	HEMBA1003725	46. 157	55. 932	71.704	30.085	21. 305	22. 378	18.643 J	31.573
	HEMBA1003728	103.795	35, 668	58. 184	16.485	21.818	42. 286	37.790	34. 280
10									
	HEMBA1003729	49. 957	21. 508	47.663	20. 231	15.376	18. 567	21.294	17. 427
	HEMBA1003732	13.069	1.953	6. 558	3. 228	2.195	3.652	3.024	4. 336
	HEMBA1003733	52.409	32.781	76.684	22.919	83.426	18. 921	13.867	14. 220
	UEUG 1 1002742	40. 426	20. 265	50, 667	26. 589	21.518	42. 057	44.130	24. 802
	HEMBA1003742								
	HEMBA1003743	26.918	22.118	23. 392	18.886	18.530	12. 506	17. 162	18.069
	HEMBA1003758	110.630	126. 359	315, 104	79. 435	58. 130	58. 587	34.868	73.429
15									
7.5	HEMBA1003760	78.949	0.000	26.318	15, 194	14. 440	32.057	34. 468	19.471
			30. 390	82,720	23.891	19.630	164. 051	37.797	57.861
	HEMBA1003764	45. 855							
	HEMBA1003769	87. 589	47.227	62. 942	27.144	32.047	46.499	39.296	38. 944
		63.842	14.722	21.132	12.002	9.850	33. 904	29, 817	13. 165
	HEMBA1003773								
	HEMBA1003783	17. 751	16. 975	23. 942	16.465	13.884	6.842	9. 757	20.650
		13.500	17, 233	21.849	13.856	12.436	17. 394	11.099	13.140
20	HEMBA1003784								
20	HEMBA1003794	386.642	303.008	322. 299	109.371	145.316	286.778	287.377	239. 938
	HEMBA1003799	39. 392	23, 099	29.603	15.022	13.775	16. 550	24.428	19. 403

	HEMBA1003803	63. 548	21.899	44. 323	20.132	18.580	28.795	24.744	35. 938
	HEMBA1003804	80.382	26, 816	48. 558	16.154	27.867	31.087	37,611	22.634

	HEMBA1003805	103.669	42. 485	42.930	19. 994	36.377	43. 797	32.147	28.376
	HEMBA 1003807	21.717	13.940	25. 512	9, 492	6.870	9. 649	8.812	7.611
25	HEMBA1003810	20. 102	11.572	7.558	20.338	17.855	7. 640	4. 451	6.585
	HEMBA1003827	432.964	219.520	240. 291	155.416	219.584	266. 037	283, 204	241.127
	HEMBA1003836	177. 311	135.831	482.334	146.466	136.063	93.790	92.728	122. 237
	HEMBA1003838	223.674	185. 295	641, 368	134.002	79.993	115.711	87.137	118.957
						21, 104	13. 382	11.701	13.634
	HEMBA1003843	13.867	10.178	27. 409	17.850				
	HEMBA1003846	133.994	57. 556	58.738	34, 962	50.550	56. 395	40.861	60.253
					14.248	8.662	11. 259	9, 145	9.934
30	HEMBA1003856	27.378	13.868	16. 982					
	HEMBA1003857	101.908	95. 527	253. 525	75.110	52.628	51. 958	45.837	48.871
				24.567	9.568	13.009	16.810	29, 271	16.795
	HEMBA1003864	52.130	18.071						
	HEMBA1003866	27.257	12.805	22, 440	12.069	15. 414	19. 103	9. 229	7.524
			54. 991	58. 923	31,090	41.733	69. 461	48.174	43.486
	HEMBA1003868	95.701							
	HEM8A1003879	62.950	44. 572	159, 217	48.098	42.446	37. 097	36.010	45. 824
	HEMBA1003880	134.462	70,074	103. 271	50.699	47.956	67.668	44. 498	30.581
35									
	HEMBA1003884	99. 190	48, 465	73.499	34.796	54.399	57. 269	63.551	68.830
	HEMBA1003885	77.675	69.096	172.968	55, 129	49. 424	41.309	24. 247	31.596
	HEMBA1003887	60. 203	22. 185	33.582	16,896	21.181	29. 281	31.275	22.835
	HEMBA 1003890	12.753	8.056	15. 506	7, 762	16.057	139. 271	387.408	5.124
							212. 964	137, 297	122. 335
	HEMBA1003893	386. 525	281.955	515.307	187, 300	180. 355			
•	HEMBA1003895	411.418	232.899	382.182	1144, 104	165.806	233. 857	186,700	143.577
40		39. 732		114.984	20.297	23.509	16.793	14, 124	20. 479
	HEMBA1003902		39.491						
	HEMBA 1003 904	32.775	21.109	45.629	10.006	13.109	14. 294	24. 342	17.444
	HEMBA1003908	8.660	8.873	15.689	7.298	15. 429	6. 307	2.267	5.699
	HEMBA 1003926	132.636	253.614	316.882	183.017	124. 195	147. 955	105, 962	360. 995
	HEMBA1003937	87.005	63, 862	200, 940	40.687	36.238	35. 284	29.695	40, 418
								10 153	15 202
	HEMBA1003939	28.064	25. 844	35.675	20.306	20.378	19.070	16.45/	15.626
45	HEMBA1003940	27, 800	13.368	18.045	10, 235	10.394	14.633	17.733	9.868
							23. 428	19.757	13.795
	HEMBA1003941	57.997	16.835	24. 582	17. 381	15.884			
	HEMBA1003942	38.168	19.747	45. 852	32.660	22.333	24. 695	10.791	21.900
							36. 632	32.279	26, 903
	HEMBA1003945	59. 457	32.900	46.079	23.037	21, 163			
	HEMBA1003949	12.870	13.019	20.678	7, 159	38. 521	442. 120	272.494	21.625
						4.756	3, 396	8.814	5, 401
	HEMBA1003950	8. 366	8. 726	5.814	3, 195				
50	HEMBA1003953	23. 527	10.310	11.872	9.390	8. 494	10.637	10.973	5. 252
			90.718		74. 499	85.036	62. 450	34. 852	86. 629
	HEMBA1003958	131.082							
	HEMBA1003959	12. 105	11.228	18, 520	6. 548	7.960	18. 122	12.612	10. 591
	HEM8A1003960	53. 133	29.785			16, 178	21.708	32.094	35. 333
	HEMBA1003966	58. 245	19.415	68.506	20.791	26. 975	28. 975	27.825	25. 303
	HEMBA1003967	1.859	3.908	9.364	6.033	4.054	4. 384	4. 208	4, 985
	LIFE CON LOCATION	1.000	3. 300	3, 557	1 000	, ,,,,,,,			
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Table 28

	Original								
	HEMBA1001968	40.219	26.894	55. 357	16.296	14.511	28. 531	22. 648	15. 420
	HEMBA1003974	147. 167	439. 547	139.030	117.010	33.973	54. 122	29. 356	338.820
_	HEMBA1003976	20. 167	17, 809	13. 159	9.187	5.748	6.820	6.962	10.367
5	HEMBA1003977	32, 751	12.350	24. 212	6. 558	6.776	12.413	17.016	9.367
i	HEMBA1003978	40. 564	13. 858	10.812	11.585	11, 203	23, 881	20. 489	
									17. 488
	HEMBA1003981	65, 803	34. 462	71.399	26.801	31.348	48.051	31.355	42.728
	HEMBA1003982	15, 104	89.360	20. 946	18.086	1.620	3. 781	3. 102	64.356
	HEMBA1003985	15, 199	10.866	21.715	9. 199	1.517	8.041	5. 977	7.569
	HEMBA1003987	48.695	30.080	108.473	25. 632	23. 222	28.008	21.302	24. 940
10	HEMBA1003989	47. 841	51.466	128.889	32.288	24. 298	24.627	15. 392	23. 174
	HEMBA1004000	36. 424	35.098	34.843	16. 292	19, 541	20.604	16. 803	21.872
	HEMBA1004006	8. 411	42. 393	12.931	2.863	3. 395	0.000	4. 943	9.742
	HEMBA1004007	135. 300	114.014	286,000	90.971	64.473	74. 153	71. 985	79.319
	HEMBA1004010	58. 331	152.845	38.786	18.676	18.819	35. 229	31.514	80.599
	HEMBA1004011	62.306	16. 294	38. 336	12.356	13.756	29. 583	26.091	7.986
15	HEMBA1004012	47.010	38. 053	139, 110	42.415	22. 159	34. 340	27. 215	32.550
	HEMBA1004015	24.416	26. 249	27.372	12.243	13.962	25. 082	25. 133	12.269
	HEMBA1004024	149. 457	114.788	479.037	80.67 9	77.896	75.066	57. 366	93.859
	HEMBA1004029	81.485	31.944	43. 520	19.897	20. 191	38.768	36. 482	19.376
	HEMBA1004038	26.629	15. 823	19.708	12.109	7.832	14. 400	12.855	17.771
	HEMBA1004042	8, 177	10.678	12.830	6.612	11.484	7.963	11.320	10.405
20	HEMBA1004045	24.675	30.855	37.128	20.069	23. 538	15, 509	17. 299	17.447
		95. 795	48. 977	78.760	36.608	40.779	45. 132	47. 334	63.844
	HEMBA1004048	35. 133							
	HEMBA1004049	55. 947	543. 954	47. 428	49.034	19.297	56. 209	23. 320	68.865
	HEMBA1004051	69.776	31.608	51.948	13.046	25.684	38. 632	30. 423	32.553
	HEMBA1004053	29. 222	70.670	84. 481	24, 394	15.007	23. 414	13. 218	23. 973
	HEMBA1004055	39.564	23. 202	34. 928	8.151	5. 353	28.619	15. 237	14.807
25	HEMBA 1004056	136. 121	122.072	413.353	75.363	81,883	56. 439	41.004	85.794
	HEMBA1004060	17.642	11.826	29.995	9.507	4.910	13.895	8. 579	8. 388
	HEMBA1004061	17, 144	13. 460	20.009	16.913	8. 228	14. 145	12. 424	5.810
				104. 390	62, 419			94.004	
	HEMBA1004067	165.029	79.589			50.783	89.115		91.850
	HEMBA1004071	28. 405	34. 722	37.707	19.775	14.692	17.342	23.864	27.554
	HEMBA1004074	128. 445	51. 388	148.050	35, 606	37.851	50.216	53. 461	46.373
30	HEMBA1004078	26. 126	14.714	20.940	9.721	16.211	17.398	17. 388	14.057
	HEMBA1004085	42.006	24. 067	36.862	15.417	17.609	19.555	28. 362	21.993
	HEMBA1004086	27. 330	49. 843	21.238	43.213	24. 232	16. 260	12.409	22. 262
	HEMBA1004097	45. 296	15. 292	27.795	13.971	26. 928	26.002	33. 192	19.361
	HEMBA1004100		37. 210	48. 942	23. 245	10. 184	25.744	21.452	28. 594
		40.930							
	HEMBA1004103	101.036	101. 281	184.668	64.176	44. 322	55. 385	41.050	40.000
35	HEMBA1004110	89, 903	65. 107	57.751	43.841	27.836	21.315	27. 631	34. 280
**	HEMBA1004111	171. 907	134. 108	296.310	95. 474	115.874	78. 450	80.011	98.760
	HEMBA1004124	177. 408	71.838	103.065	37.865	46. 198	68. 531	109.364	77.083
	HEMBA1004130	64. 543	54. 797	171.602	50.628	35. 382	25, 601	19. 599	23.097
	HEMBA1004131	41.654	24. 184	33. 975	26.913	23. 365	28.790	20.022	24. 999
	HEMBA1004132	55. 906	42.840	162. 243	42.708	30, 251	28. 863	19.780	22.237
	HEMBA1004133		30.838	38. 522	29.390	20. 897	28. 027	28.747	33. 333
40		64.624							
	HEMBA1004138	61.197	21.853	23.858	17.376	9. 337	30.080	17. 345	22.082
	HEMBA1004143	15.715	9. 656	21.209	10.565	10.539	14.067	11.441	9. 994
	HEMBA1004146	40.893	21.789	90. 537	30.633	32.870	23. 542	14. 368	20. 982
	HEMBA1004148	59. 990	18.796	22. 167	11.049	17.531	18.309	29. 374	22.628
	HEMBA1004149	16, 284	11.131	18. 385	7.758	7.634	7.577	5.890	13.683
	HEMBA1004150	5. 223	4.403	4.468	3.044	2.553	2.158	2.062	2. 260
45	HEMBA1004154	111.110	40. 836	69. 965	31.437	46. 253	58. 472	62.983	47.866
• •	HENBA1004164					47.327			
		139.670	107. 565	315. 189	77. 326		57.372	46.726	67. 257
	HEMBA1004168	24. 042	16. 530	18.698	9.347	9.400	13.838	3.054	13.060
	HEMBA1004199	22.894	9. 047	10.461	8.631	7.704	7. 849	6.889	7.253
	HEMBA1004200	33. 301	51.362	83.462	26. 185	27.548	17.580	17. 235	32, 109
	HEMBA1004201	54.766	23.783	32.370	17.449	21.835	22.123	25.993	20.006
50	HEMBA1004202	14. 526	10.484	12.784	6.804	5.704	9. 594	8.672	11.673
	HEMBA1004203				13, 604	14. 171	19, 946	16.079	
	III. III. III. III. III. III. III. III	47.655	20.140	34.882					18. 151
	HEMBA 1004207	6.344	3, 206	11.421	3.936	6. 145	5. 704	21.692	7.780
	HEMBA1004210	33.071	43. 543	33.120	16, 340	41.396	21.814	19.639	15.015
	HEMBA1004225	73. 182	63.749	226. 133	59.565	43.156	32.703	25. 781	40.078
	HEMBA1004227	83.820	31.222	42.541	16.931	17.786	28. 177	25. 468	30.978
55		1 -3. 3-9			•				

. Table 29

	HEMBA1004235	99. 954	57.144	62.536	27.672	34. 345	69.613	47. 182	38.807
	HEMBA1004237	27.504	21.542	17.029	18. 289	11.697	19.212	12.031	16. 922
					34. 130	27.841	36, 089	27.438	34. 578
5	HEMBA1004238	79.210	38. 454	102.493					
	HEMBA1004241	5.663	2.654	7. 035	2. 556	1.072	2. 912	4. 422	1. 294
	HEMBA1004242	256.862	65.757	191.327	80.010	76.455	85. 478	89.242	62.567
	HEMBA1004243	72.699	55. 276	60, 764	28. 287	47.148	36.800	28. 491	47.743
	HEMBA1004246	44.915	30. 967	100. 300	22.414	17. 109	15. 470	12.686	18. 700
	HEMBA1004247	66.750	16. 238	24. 674	18.889	22.763	31.897	38.415	17. 377
	HEMBA1004248	13.953	18. 412	17. 581	11.953	11.378	14, 538	12.794	9. 562
10	HEMBA1004250	24. 439	10.494	10, 631	6.401	5.142	14. 218	12.652	11.966
	HEMBA1004252	37. 349	20.650	22. 246	9. 949	9. 550	14.570	21.841	18, 200
	HEMBA1004260	10.994	19.320	16.415	15.707	20.374	13.845	11.265	19. 838
	HEMBA1004264	22.716	14.715	13, 358	7.615	5. 234	12. 282	15.089	11.397
		235. 310	195.750	654. 331	171.071	174. 292	115.073	102.973	144. 125
	HEMBA1004267								
15	HEMBA1004272	28.776	19.025	23. 678	13.063	12.012	15. 529	14.123	14. 593
10	HEMBA1004274	62. 157	50. 491	53, 598	30.356	36.472	42.005	58,020	, 51. 617
	HEMBA1004275	70. 423	38. 514	45, 176	17.443	18. 132	34.031	36.295	22. 171
	HEMBA1004276	13.630	4. 481	14.011	9. 548	9.099	14.035	10.406	8. 615
	HEMBA1004279	16.536	11.082	13.356	14.834	7. 333	10.255	8.919	12.068
	HEMBA1004284	29.688	30. 297	64.483	13.658	17.646	17.327	17.630	13.770
	HEMBA1004286	32. 471	16. 566	18, 049	12.391	6.773	17.625	23.811	13. 547
20	HEMBA1004289	81.573	62.930	165, 571	49.704	34.785	37.379	28.939	41.740
							57.764		
	HEMBA1004293	72.456	34. 902	48, 669	32.705	17. 408		53.695	45.065
	HEMBA1004295	37. 595	12.116	29. 975	11.634	5. 514	25.018	23.797	20. 926
	HEMBA1004302	10.880	5.912	7.885	10.025	5.190	6.060	5. 264	9. 355
	HEMBA1004306	426.811	177. 321	335. 158	107.645	123. 947	256.397	251.772	134. 005
	HEMBA1004312	37. 953	30.864	105. 533	30.747	25.847	16.140	16.283	24, 272
25	HEMBA1004314	29.396	23.332	95. 584	22. 179	18.544	11.015	8,804	20. 974
	HEMBA1004321	47.670	29. 150	105.316	35.655	23.139	31.309	29.736	47. 858
	HEMBA1004323	87. 295	65. 931	221.440	44.690	41.425	36.609	34, 117	39.135
	HEMBA1004327	65.869	21.284	21.540	11.985	14.419	27.213	27.030	20.118
				132, 755		21.556	32, 356	17.478	26.773
	HEMBA1004329	67. 920	44. 687		32.977				
	HEMBA1004330	8.765	7.655	16.827	7.164	3.843	9,511	7.660	4.615
30	HEMBA1004334	16.438	21.355	31.680	15.109	26.670	13.368	10.581	13. 568
	HEMBA1004335	204.961	102.859	325. 226	69.979	64, 392	78.772	71.641	83. 525
					15. 995	41.404	87.221	89.558	40. 224
	HEMBA1004341	186.677	30, 208	61.439					
	HEMBA1004344	261.676	76.316	123. 332	42.705	51.432	26.797	42.054	59.071
	HEMBA1004347	65. 249	32.610	97.858	37.038	21.953	33.115	33.526	36.846
	HEMBA1004349	22. 353	35. 727	29, 441	19.803	18.786	23.126	19, 103	18.719
	HEMBA1004352	75. 508	65. 544	237.050	49. 039	34. 141	32.597	28.166	46.343
35						27. 380	39.551	30.556	56.886
	HEMBA1004353	54. 322	66.042	132.169	40.563				
	HEMBA1004354	43.687	29. 352	79. 264	22.784	20, 533	21.755	16.860	22.429
	HEMBA1004356	44.730	22. 201	27.487	10.404	8.280	22.159	16.039	15.038
	HENBA1004360	91.412	28. 429	71.634	26. 232	36, 259	59.602	38.361	50.410
	HEMBA1004366	9. 956	10.099	14. 263	5. 481	5.631	6.802	6,791	6. 167
							1.507	3. 555	
40	HEMBA1004372	3,613	4, 593	5. 338	0.000	1,638			1.568
	HEMBA1004377	53.834	41.410	47.048	29. 140	26. 163	34. 545	30.827	33. 572
	HEMBA1004389	20.540	22.800	24.474	14.497	13.968	16.620	14. 951	17.114
	HEMBA1004391	60.284	22, 653	44.013	14. 283	19.018	31.716	23.931	23.617
	HEMBA1004393	177, 786	197.548	108. 554	32. 455	75.399	76.587	39.777	44.665
	HEMBA1004394	28, 949	11.849	12.442	5.544	10.440	17. 825	10.981	8. 836
	HEMBA1004396	37. 907	26.956	102.760	18. 571	16.519	15.025	13.681	21. 980
45	HEMBA1004401	22.519	21.858	30. 601	14.945	13.592	15.418	20.530	20.774
	HEMBA1004405	42.933	38.835	117.844	34. 528	23.557	19.155	18.506	29.842
						15. 351	19. 522	15.546	20.863
	HEMBA1004408	50.497	27. 151	55.000					
	HEMBA1004414	45, 769	51.722	64.316		19.324	39.735	26.527	36. 385
	HEMBA1004429	61.867	59.067	190.058	39.014	50.304	38.462	27.517	46. 317
	HEMBA1004433	49, 568	39.828	146. 938		28. 383	24. 241	24.651	42.005
50							23. 183	21.969	35. 073
50	HEMBA1004440	31.849	22.499	37. 132		22.366			
	HEMBA1004444	59, 488	46.586	163.763		29.990	24.833	22.908	37.635
	HEMBA1004446	22, 134	12.309	29. 426	11.920	3.385	14.862	10.855	16.078
	HEMBA1004451	31,688	21.261	28.136		15.678	19.695	21.159	27. 156
	HEMBA1004452	36. 593	5. 268	18. 479		5. 737	17.680	14. 173	7.972
	HEMBA1004454	50.056	27. 897	32.786	24. 382	20.631	24. 494	22.897	29. 042
55									

Table 30

•	HEMBA1004460 T	120 CEN T	96, 143	356.058	74. 883	54, 735	70.698	38.344	55.945
	HEMBA1004461	138.550 64.074	19.163	24. 808	12.846	16. 373	23. 508	22.827	10, 137
	HEMBA1004468	134. 439		210. 409	77.409	60, 142	56.229	42. 361	49, 457
	HEMBA1004479	82.994	32.899	92. 282	35. 519	44. 435	43. 183	31.836	37, 116
	HEMBA1004482	5, 602	7.682	11.248	36.034	2. 926	5. 535	5.693	5, 972
	HEMBA 004491	16, 736	6. 285	17.615	14,018	11.729	19.804	10.683	11.280
	HEMBA 004499	94, 095	71.456	148. 355	58, 479	48, 596	46. 968	46.648	57.279
	HEMBA1004502	21. 523	9. 344	18.265	7.282	9, 979	2.762	3.174	13.389
	HEMBA1004505	26.042	15. 980	43. 855	17.516	15. 469	22. 190	22.873	15.812
	HEMBA1004506	12.004	29.395	42.664	21.849	21.426	14.469	9. 224	9.845
	HEMBA1004507	96. 377	87.688	99. 177	03.472	34, 160	81.068	54. 939	151.142
	HEMBA1004509	52.657	14.880	19. 120	16. 228	17.009	24.783	24. 565	13.476
	HEMBA1004523	20. 156	18.209	11.197	16.529	14, 651	13.004	20. 267	19.467
	HEMBA1004528	42.620	27.819	48.069	14. 426	25. 267	43.038	40. 239	37.718
	HEMBA1004534	75.090	41.159	44. 399	31, 300	16.686	31.317	21.009	18.589
	HEMBA1004536	31.531	13.343	23.664	14.085	4. 408	13, 133	12. 981	15. 201
	HEMBA1004538	352.363	181.508	233.819	97.018	122. 402	183.507	100. 197	150.062
	HEMBA1004542	47, 360	17.733	29. 238	17.280	12.324	17. 317	22.764	15. 212
	HEMBA1004552	63.401	29. 585,	26.857	43. 567 16. 273	39.674 32.820	38.686 33.809	23.830 20.812	33. 542 5. 536
	HEMBA1004554 HEMBA1004558	62. 231 30. 217	11.953 12.133	25. 084 31. 036	15. 840	30.638	65. 183	19. 155	30. 921
	HEMBA1004560	68. 901	16.556	17. 908	16. 431	10, 034	30.093	22.379	22.683
	HEMBA1004564	48. 119	14. 911	35. 565	31.983	32.464	30.028	20. 965	32,479
	HEMBA1004566	32, 479	29. 553	20.970	32.788	42.949	40.715	23. 273	32.960
	HEMBA1004573	17.728	13.843	7.118	9.972	19, 952	9.755	9. 278	8.100
	HEMBA1004576	39. 572	26.733	42.044	10.704	37. 505	32.441	17. 232	25.361
	HEMBA1004577	46.233	11.570	97.881	39.434	13. 437	41.089	34.426	35. 314
	HEMBA1004586	82.532	71.398	213.814	70.289	52. 589	45.729	23. 395	38.312
	HEMBA1004596	72.534	32.493	45.820	27. 585	27.854	34. 997	33.847	38. 473
	HEMBA1004604	99.019	48. 582	103. 587	36.723	49. 392	48. 377	56. 558	69.256
	HEMBA1004607	53.557	37.013	100. 999	27.559	26. 143	28.796	21.692	42.044
	HEMBA1004610	20.690	14.854	69. 908	15. 349 22. 819	12. 120 15. 568	9, 108 10, 691	8. 858 6. 697	15.087
	HEMBA1004617	22. 592	20. 386	42. 426 209. 059	49. 931	29, 836	29. 902	12. 194	27. 438
	HEMBA1004622 HEMBA1004626	78. 025 38. 170	46.803 36.312	110.684	22.791	14.118	17, 193	15. 579	20.821
	HEMBA1004629	33.858	37. 886	87. 440	53, 228	47. 341	28. 160	12.170	28.096
	HEMBA1004631	35. 946	10.475	4. 434	7.390	17, 128	22.775	9. 569	32.852
	HEMBA1004632	27.084	13.891	23.598	10.209	7.802	11.754	22.566	6.362
	HEMBA1004633	78.391	33.135	114.054	17.197	49.008	60.659	48.857	40.810
	HEMBA1004636	52.397	20.706	34.962	10,085	22.609	21. 255	13.502	25.039
	HEMBA1004637	4. 228	4.304	6.747	5. 278	9.756	4.086	2.597	5.024
	HEMBA1004638	0.241	0.000	0.000	1.008	0.000	0.000	0.113	0.000
	HEMBA1004645	57.971	29. 263	111.067	32.645	17. 998	27. 214	20. 560	24. 845
	HEMBA1004656	16.139	9.194	21.399	12.766	18.216	14.099	17. 122	12.004
	HEMBA 1004657 HEMBA 1004666	20.820	23.742 3.174	69.842 18.097	9, 422 5, 962	138. 932 9. 830	42.697 5.098	9. 048 2. 525	13. 383 7. 512
	HEMBA1004669	7.321 94.910	36. 291	111.210	30.591	20.021	28.018	25. 500	25. 624
	HEMBA1004670	57.231	17.070	60. 538	23.280	13.173	24. 312	23.413	14. 342
	HEMBA1004672	63.471	50.154	146.619	39.883	31.559	25.617	20.328	28.099
	HEMBA1004689	152.993	93. 435	103.311	81.212	50.901	83. 998	57. 329	84. 276
	HEMBA1004690	28.240	10.247	13.401	8.159	4. 952	13.963	13.991	11.785
	HEMBA1004693	18.359	15. 228	20.803	14.290	13.070	16.726	9.014	13.531
	HEMBA1004697	81.532	48.847	148. 587	58.849	34.416	51.983	42.641	50. 271
	HEMBA 1004702	97.518	62.966	49. 904	20.714	42.224	58. 936	64. 906	37. 506
	HEMBA1004704	99.561	48.717	236. 687	38.866	33. 457	38.377 10.638	24.626	31.783 4.810
	HEMBA 1004705	12.717	12.313	40, 950 16, 175	9.649	17.803	17. 906	5. 969 13. 036	12.703
	HEMBA1004706 HEMBA1004709	33.616 51.126	9.825 39.934	136. 723	32.285	25. 072	21.674	15. 230	23.755
	HEMBA1004709		9. 203	57.020	12.805	14. 304	16.154	12. 982	9, 790
	HEMBA1004711	121.283	47. 643	73. 497	30.236	56.917	65.719	56. 298	52.009
	HEMBA1004725	56.905	32.051	70. 171	12.221	48.208	34.021	35. 739	12.501
	HEMBA1004730	36.072	10.037	30.016	7.633	13.361	7.545	8. 989	34. 832
	HEMBA1004733	30.769	29.884	23. 348	6.988	2.998	8.055	8.031	2.822
	HEMBA1004734	11.912	11.974	36.595	3.988	12.556	7.653	4. 303	15.670

Table 31

	HEMBA1004736	55. 309	25. 331	132.333	45.653	38.696	23.516	19.970	34. 509
	HEMBA1004748	53.832	20.004	172.069	29. 562	22. 161	14.904	12.665	18. 349
5	HEMBA1004749	127. 285	45. 137	73.698	27.788	33. 184	60.214	44.636	42.250
-	HEMBA1004751	81.283	64.830	173.888	54. 165	36.368	41.802	31.283	43.505
						29.090		30.970	33.029
	HEMBA1004752	59.058	32. 785	109. 428	32, 254		34. 259		
	HEMBA1004753	204.044	247.466	406.165	156.689	102.755	131.323	83.515	263.882
	HEMBA1004755		59.677	83.850	22.148	29.800	30.642	13.064	23. 261
		57.638							
	HEMBA1004756	9. 965	16. 228	11.023	8.349	5.780	9. 109	111.528	14. 885
	HEMBA1004758	36.487	26.558	116.970	22. 341	14. 553	14.773	11.840	14.406
10							25. 489	23.061	
	HEMBA1004763	67.343	19.641	33.742	13.841	16.720			18.650
	HEMBA1004768	29. 177	24. 043	38. 303	6.673	10.298	3. 197	10.352	13. 391
	HEMBA1004770	10. 327	14. 492	10.901	6.416	6.310	7.963	10.868	7.955
	HEMBA1004771	46. 910	34. 314	76.491	31.509	22.830	23. 102	30. 433	32. 358
	HEMBA1004775	39. 253	28. 706	63.968	24. 931	18.754	43.049	32.720	26.795
						9.000	16.400	10.105	8.046
15	HEMBA1004776	22.604	11.017	10, 103	5. 466				
,,,	HEMBA1004778	78.144	77.681	223. 475	37.540	33.791	32. 337	24.057	43.529
	HEMBA1004784	9, 826	18. 370	102.812	8.313	15. 151	11. 373	9.479	6.329
	HEMBA1004785	25. 723	16. 345	26.216	6.651	10.649	10.674	13.732	11.615
	HEMBA1004789	18. 173	14. 508	16.096	7.804	8. 691	10.011	7.713	11.389
							9.757	9. 905	12.028
	HEMBA1004795	14, 283	12. 973	25. 122	11.028	9. 351			
	HEMBA1004797	65. 927	33.745	73.888	34. 142	28. 246	40.067	32.715	25. 583
20	HEMBA1004803	36.634	41.124	65.880	27.072	30.957	22.607	22.520	26.554
						9.653	9.000		8.399
	HEMBA 1004806	11. 997	8. 183	21.467	8.868			7.894	
	HEMBA1004807	16. 352	14.481	22.459	11. 249	12.009	13.340	7. 935	9.118
	HEMBA1004816	29. 782	24.075	95.884	18.110	29.259	8.180	12.578	10.934
	HEMBA1004820	8. 636	7.466	8.862	4. 249	4.018	4. 269	6.876	3. 493
	HEMBA1004833	159. 947	50.729	81.248	38.650	64.754	83.155	56.657	65. 121
25	HEMBA1004847	51.456	25. 570	40.694	21.115	36.572	35.053	31.537	40. 529
20									
	HEMBA1004850	77. 254	24.014	38.620	21.854	26.080	54, 413	50.197	24, 185
	HEMBA1004863	57, 117	32.704	72.480	23.951	31.887	25.058	20.050	20. 982
		46.043	27.344	59.824	26.750	13.898	16.719	20.308	17.843
	HEMBA1004864								
	HEMBA1004865	12. 257	14.642	31.748	44.090	14. 331	13.454	13.835	15.797
	HEMBA1004880	56.788	50.021	126.837	35. 420	26. 589	24.064	20.647	23. 264
				29.340	16.782	13.013	13.652	10.676	19.977
30	HEMBA1004882	42.450	18. 453						
	HEMBA 1004885	8. 545	4. 947	5.350	4.891	2. 933	3.711	3.652	6.615
	HEMBA1004889	28. 103	22.485	32.049	17.078	14. 363	23.391	15.605	16.916
							6. 539	5. 245	9.440
	HEMBA1004900	19. 922	15. 709	33.254	10. 423	9.045			
	HEMBA1004909	88. 522	49. 269	163.284	48. 147	35. 537	36.045	18.861	27.933
	HEMBA1004918	64. 384	43.134	105.868	34.899	22. 323	24.073	15.857	25. 370
35	HEMBA1004923	47. 731	37.996	69.168	19.659	26. 441	18.192	10.213	20.111
33	HEMBA1004929	11.048	14.003	10.808	12.050	7.539	9.882	8.967	11.809
	HEMBA1004930	101.277	92.425	279.652	80.664	66.618	34. 331	31.091	41.874
	HEMBA1004933	9. 145	5. 566	12.895	7.786	12.296	10.327	96.467	5. 417
	HEMBA1004934	7.311	7.106	43.966	10.208	4.750	5. 866	9.143	12.805
	HEMBA1004937	43. 331	27.219	38, 802	15. 368	17, 734	15. 280	15.784	46.365
40	HEMBA1004943	51.072	26.833	32.001	21.614	16.458	27. 585	29.628	38. 533
40	HEMBA1004944	84. 363	46.788	126.294	43.803	28. 989	38. 514	31.589	23.074
	HEMBA1004946	64.638	28. 144	37.908	17.163	24. 332	27.854	34.636	31.712
	CEMON 1004340	NA 222					22 550	20 868	
	HEMBA1004952	90.835	18.893	40.862	12.824	20.090	33. 568	20.062	19.020
	HEMBA1004954	14. 656	36.003	41.485	27. 126	23.696	20.777	6. 946	29. 261
	HEMBA1004956	5. 975	9. 923	6.635	7.743	0.953	4.578	1.565	5.188
	NEMO/1004330								
	HEMBA1004960	86.030	77.420	136.061	60.735	49. 221	47.560		45. 929
45	HEMBA1004971	31.046	5, 439	7.559	12.468	17.946	15.068	19.705	18.480
-						27. 295		30.004	50.710
	HEMBA1004972	77. 318							
	HEMBA1004973	35. 524	13.502	16.731	9. 641	11.726	14.716	19.197	22. 580
	HEMBA1004977	6.756	9, 870			29.373	8.701	2.217	10.523
	HEMBA1004978	8. 589				5, 158		2.642	10.106
	HEMBA1004980	34.093	33. 440	87.268	25, 974	18.071	16.453	11.605	22.124
E0.	HEMBA1004982	14.750		17. 944		8. 250		6.083	5. 456
50									
	HEMBA1004983	38. 285	13. 488	20.831	11.831	3.348		11.455	8.305
	HEMBA1004995	27. 256	28. 515	26.297	18.434	25. 474	22.491	24. 452	33.683
						7.816		7.576	14.698
	HEMBA 1005004	13.855							
	HEMBA1005008	64.714	26.633	22. 502	18.478	23. 532	28.617	18.581	16. 940
	HEMBA1005009	34. 543				14, 122	26. 432	12, 593	23.116
	[:::::::::::::::::::::::::::::::::::::	1 07. 373			1 . 5 . 5 . 10				
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Table 32

	HEMBA1005019	49. 260	24.872	25. 349	20.834	30.144	32.629	20.777	27.016
	HEMBA1005021	37.224	63.713	38.065	17.061	15. 647	20.369	25. 859	32.656
5	HEMBA1005029	30.265	17.783	35. 352	16.531	19. 588	26. 517	15.798	16.604
•	HEMBA1005035	393. 404	200.167	574.746	133.872	210.689	177.872	156.563	170.510
	HEMBA1005036	115. 345	41.961	73.015	39. 541	44. 451	66.623	55.833	51. 349
	HEMBA1005039	28.850	19.922	57.018	13.971	11.999	26.427	15. 206	7. 350
	HEMBA1005047	93.995	31.868	54. 335	18. 576	28, 338	31.562	31, 930	23.751
	HEMBA1005050	78.015	41.690	73. 330	29.830	26. 504	35. 887	21.640	35, 653
10	HEMBA1005062	23.050	15.803	29. 553	15.707	7.836	15.618	19. 435	13. 336
10	HEMBA1005066	10.980	11.364	31.553	13.509	5. 668	10.541	5.005	10.849
	HEMBA1005067	39. 308	34. 578	39.795	44.519	24. 643	21.272	19.379	20. 121
	HEMBA1005070	73. 155	34, 949	68.556	29.956	38.004	38. 211	48. 007	31.733
	HEMBA1005075	88.089	37.798	148.675	40, 537	33. 271	33.074	28. 661	30. 201
	HEMBA1005078	100.064	37.746	66.827	33, 115	41.170	55. 560	51. 231	17. 112
40	HEMBA1005079	137.757	86.238	294.118	73.304	76.035	75.084	47.255	76. 170
15	HEMBA1005083	18. 102	7.642	17.087	6.711	6.184	8. 675	9. 287	9, 609
	HEMBA1005084	82.712	38. 248	47.063	26. 564	27.435	37. 552	38.419	28. 349
	HEMBA1005088	31.610	22. 435	76.774	22.700	18. 926	23.875	8.895	12.447
	HEMBA1005089	68. 944	55. 156,	178, 226	34.742	32.350	38. 645	22.869	28.148
	HEMBA1005090	148.861	86.156	117.997	94.811	57.034	81.098	54. 187	116.066
	HEMBA1005096	83.125	30.911	63.940	33. 378	33.962	48. 589	35.467	36. 021
20	HEMBA1005101	69.080	14. 020	34. 136	10. 197	13.998	34. 420	22.696	15. 975
	HEMBA1005107	82.659	25. 203	36, 223	11.215	21.514	32.720	25, 972	21.337
	HEMBA1005113	7.977	17. 225	31. 501	7. 563	44. 493	5. 157	6. 957	9.761
					7.303				
	HEMBA1005123	173.637	77. 260	555, 672	126.908	94.628	90.446	70.735	90.016
	HEMBA1005133	58. 192	40, 749	122.920	29.864	16.700	17.652	8.802	18.988
	HEMBA1005135	8. 259	9.125	14.962	2.213	16.732	6.892	3. 383	6.189
							0.692		
25	HEMBA1005145	185. 299	101.220_	352.159	92.082	88.750	122.118	76.475	90.044
	(HEMBA1005149	220. 122	109.352	274.492	120.663	125. 192	96.704	92.083	128.030
	HEMBA1005152	125. 948	96.291	226.882	58. 505	33.738	46. 323	27.534	34. 457
	HEMBA1005159		11.274	9. 399				12.001	
		15.760			6. 198	6. 191	6.861		4. 556
	HEMBA1005172	1653. 208	89.658	73.666	54.667	33.118	55.680	32.520	70.907
	HEMBA1005185	9.954	17, 248	10.492	34. 452	3. 558	3.117	6.026	11.173
	HEMBA1005186	23.745	10.048	27.091	13,067	7.719	15. 412	15.086	15.591
30									
	HEMBA1005195	14. 573	8.648	11.038	19.306	6.313	25. 313	13.510	9. 183
	HEMBA1005201	52. 322	13.197	47.505	13.091	12.078	8. 531	23. 532	9.848
	HEMBA1005202	98. 566	30.141	71.588	27.954	44. 381	66.294	42.390	39.695
	HEMBA1005204	184. 429	287.156	382.039	168.753		222.970	143.609	358.646
						203. 458			
	HEMBA1005206	148. 946	61.309	84.791	34. 139	49.115	66. 295	83.608	76.159
	HEMBA1005219	21.685	17.755	9, 606	8. 236	8.038	13.031	7.751	11.441
35	HEMBA1005223	79.969	42.665	113.460	40. 547	32.099	53.017	26.025	32.004
	HEMBA1005229	00 010							
	TIEMBA 1003223	26.819	9. 926	21.841	3. 135	5. 090	6.656	4.681	7.079
	HEMBA1005230	71.184	67.313	201.065	79. 279	59.679	77.484	47.808	66.511
	HEMBA1005232	7.374	6.386	17.522	8. 552	3. 285	12.098	4. 975	3.965
	HEMBA1005238	96.780	44.134	51.932	8. 128	20.776	69. 291	49.474	35.019
40	HEMBA1005241	142. 598	104. 185	428.635	78.773	78.033	74. 434	42. 333	63.097
40	HEMBA1005244	76.771	32.597	37.797	16. 459	12.489	35. 934	31.814	35. 602
	HEMBA1005246	241.316	60.348	73.077	25.067	41.351	117.666	88. 193	54.014
	HEMBA1005251	37.505	33.247	108, 631	23. 585	14.915	23. 393	14. 302	16.409
	HEMBA1005252	53. 401	25. 532	37.199	15.002	20.744	31.279	24. 207	27. 562
	HEMBA1005267	17, 238	39, 564	20.097	27. 506	31.874	11.013	14, 526	14.024
	HEMBA1005274	16.538	8.744	18. 308	9.021	10.103	11.943	8.914	11.978
45	HEMBA1005275	69.133	43.329	216.468	46. 290	57.647	37.411	25.040	41.913
	HEM8A1005288	65.401	50.495	150.714	33.833	34.633	28. 241	24.910	40.164
	HEMBA 1005293	17.403	9, 430	23, 201	4.467	3. 192	25. 620	6.775	8.771
	HENBA1005296	223.097		894.835	738. 361	220. 523	698.319		1376.785
			811.623						
	HEMBA1005301	36.708	16.970	29.798	11.929	12.544	22. 221	35.726	32.270
	HEMBA1005304	83.978	71,914	260.016	50.686	36.101	36. 160	24.896	47, 838
50	HEMBA1005305	44. 218	33.773	74. 215	27. 494	27.352	34. 920	21.424	38.882
50									
	HEMBA1005311	33.034	20.140	48. 263	13.836	7. 908	8. 958	9.090	10.440
	HEMBA1005313	11.165	36, 175	17.550	7.047	11.502	14. 209	63.072	9.124
	HEMBA1005314	6.948	2, 955	22.604	6. 336	3.812	5.787	4.069	4.891
	HEMBA1005315								
		72.349	54.139	156.842	34. 545	43.132	26. 415	28. 942	28.442
				. 17 956	. 0 700	1 11 664	. 0 076	. 0 000	
	HEMBA1005317	20. 230	8. 451	17. 258	9.796	11.664	9. 976	9. 263	8.017
E E	HEMBA1005317	20. 230	8.451	11.238	1 9.796	11.004	9. 970	9. 203	8.017
55	HEMBA1005317	20. 230	8.451	11.238	1 9.796	1 11.004	9. 910	9. 203	8.011

Table 33

								- 277 T	7 0 10 1
	HEMBA1005318	14.755	5. 931	13.883	5. 228	5. 376	9.013	5. 511	4. 846
	HENBA1005324	98.070	33.348	44. 270	26.052	35. 446	48. 523	30, 889	17.915
								16. 942	29. 580
5	HEMBA1005331	24.826	335. 211	15. 947	25. 496	14.744	21.427		
J	HEMBA1005337	19.080	18.022	19. 429	5. 217	20.830	29. 867	32. 481	44. 585
	HEMBA1005338	61.533	38.788	63.113	23.657	30. 437	48, 455	40. 921	36.285
	HEMBA1005344	384. 481	88. 937	143.574	53.983	72.524	167.620	135. 992	68.042
	HEMBA1005353	111.629	68. 949	220.401	62.090	53.484	67.048	30, 456	42.612
				175. 543	59.707	36.743	34. 233	21.666	47. 596
	HEMBA1005359	87.635	64. 332						
	HEMBA1005362	25. 674	25.093	18.642	30.797	21.917	19.092	20. 883	12.720
10	HEMBA1005364	6.677	2.817	5, 168	13.116	19.753	5. 180	2.877	7. 198
				74, 559	28, 445	30, 138	27. 987	16.766	22.415
	HEMBA1005367	51. 911	28. 536						
	HEMBA1005372	11. 289	6.819	11.700	5.659	9. 177	6.402	9. 312	4.913
	HEMBA1005374	64.639	57. 505	120, 218	32.738	30. 987	24.792	23.695	30.728
						7.648	11.978	9. 051	4.019
	HEMBA1005379	29. 549	13.813	12.040	8.862				
	HEMBA1005382	140, 116	94.743	104, 609	70.213	26. 226	53.452	88. 235	85. 480
15	HEMBA1005384	33. 109	15, 221	21,713	10.250	8. 543	11.030	7.498	9.010
	HEMBA1005386	111.062	30.547	52.790	29.541	31.691	44.619	35. 179	29. 136
	HEMBA1005389	66.821	32.429	129, 272	42.528	35. 894	16.765	11.513	24.601
	HEMBA1005394	35.794	18. 327,	22.715	25.833	26.639	30.857	16. 944	24.443
	HEMBA1005403	40.404	14.030	54. 041	14.621	15. 504	27. 461	15. 586	32.390
	HEMBA 1005408	51.701	45.069	71.813	44. 257	67.383	35.010	23.690	44.612
20	HEMBA1005410	4.534	4. 269	11.774	12.035	10. 197	6.188	3, 955	8.910
20				163.001		50.499	41.243	22.652	35.008
	HEMBA1005411	75. 220	94.039		67.133				
	HENBA1005423	35.745	26.430	69.138	35.773	15. 442	19.286	14.057	23.010
	HEMBA1005426	14. 366	12.073	14.418	5.345	11.591	8. 954	3. 082	7. 203
	HEMBA1005427		99. 596	61.088	47.865	59.821	53.861	25. 223	46.397
		66.444							
	HEMBA1005430	52.945	15. 385	36.316	19.210	23.854	37.895	19.556	18. 127
25	HEMBA1005438	51.806	28. 359	33, 314	17.787	19. 295	21.754	13.422	29. 941
20	HEMBA1005443	108.954	165.667	426. 408	91.550	77.559	76.024	105.042	108.232
						28.000	21.457	18.763	23.755
	HEMBA1005447	51.383	39, 578	65. 244	29, 171				
	HEMBA1005449	86.452	20. 253	41.861	15.939	27.647	39. 311	28. 567	27. 508
	HEMBA1005452	110.567	52, 128	74, 119	42.532	39. 847	53. 326	67. 529	72.233
			16.821	17. 998	14.293	14. 436	8. 454	6.498	11.445
	HENBA1005454	7.997							
30	HEMBA1005468	185.066	78.008	126. 372	56.026	56. 490	78. 922	61.083	57.511
30	HEMBA1005469	88. 419	54.761	196. 280	63.682	53, 661	42.639	23.441	30.144
	HEMBA1005472	37.878	41.710	88.807	34.196	28, 126	21.983	24. 350	30. 575
	HEMBA1005474	89.169	55. 263	212.086	51.664	50.480	66. 508	39. 590	30. 322
	HEMBA1005475	212.273	98.359	182.707	110.945	105. 968	98.316	56.095	68.647
	HEMBA1005489	61.603	40.439	42, 459	21.361	21.335	31.130	11.578	25.898
					8.252	5.611	9. 949	2.213	16.039
0.5	HEMBA1005497	10.325	12.396	5. 705					
35	HEMBA1005500	86.636	39.755	180.843	46.031	28. 554	31.809	14, 951	31. 189
	HEMBA 1005506	24.029	3, 468	17. 794	7,400	5. 464	9.419	6. 593	3.517
	HEMBA1005508	12.944	12.524	22. 247	8.536	16.857	11.561	7.741	12.059
								30.786	
	HEMBA1005511	116.338	59. 193	267.636	59.921	58. 995	43.190		51.049
	HENBA1005513	167. 332	70.217	88.519	56.620	54. 920	73.797	80.751	68.624
	HEMBA1005517	37.667	10.443	23. 901	8.903	17.777	21.966	23.844	18.611
40			25. 679	71. 345	23.319	36.856	47.397	27.618	27.825
40	HEMBA1005518	109, 105							
	HEMBA1005520	200. 267	104.176	459. 373	133. 255	106. 207	95.070	67.199	94.086
	HEMBA1005522	36. 421	15, 946	24. 796	12.598	8.472	14.558	16.899	13.857
	HEMBA1005526	116. 274	72.899	292. 397	82.002	73.603	66. 198	34.319	47.682
	HEMBA1005528	13.037	9.406	30. 550	14.612	15. 947	16.518	7.583	24.988
	HENBA1005530	56.516	26. 583	63.811	13.686	21.441	29. 159	24.254	21.717
	HEMBA1005538	5.523	17.373	36. 952	7.017	10.885	11,406	15.411	35.789
45							35. 829	28.772	25. 913
	HEMBA1005539	76.498		69. 424	17.584	24. 989			
	HEMBA1005545	46. 912	10.940	32. 124	15.206	46.822	33. 595	31.865	24.090
	HEMBA1005548	57.779	14. 326	15. 050	10.139	18.638	22. 115	22. 271	39, 291
						81.893	79. 223	60.281	62.088
	HENBA1005552	141.489	120, 695	363. 831	84.934				
	HEMBA1005558	52. 488	20.021	24. 397	9.638	22.919	24.422	21.465	8.178
	HEMBA1005568	74. 152	61, 206	184. 989	53.681	38.261	33.077	24.038	37.014
50				74.768		26.562	31.212	27. 080	30. 221
	HEMBA1005570	54. 151	68.747						
	HEMBA1005576	71.454	57.260	39.016	21.283	8.931	30.461	29.371	19.991
	HEMBA1005577	40.771	13, 448	21. 181	13.021	6, 610	18.266	12.838	10.181
						19.565	33.479	28.804	16.842
	HEMBA1005581	81.577							
	HEMBA1005582	24. 681	30. 135	30. 933	14. 220	7.764	10.454	12.847	13. 157
									

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Table 34

	DELLA CACCAS I	20 504 1	66 100	00.000	0 725	10 545	10 400 1	10 533 3	17 004
	HEMBA1005583	23. 564	22.466	98.629	9.735	10.545	12.468	10.523	17. 884
	HEMBA1005588	96.041	96.264	266.022	69.126	54. 588	44. 105	34.310	52. 441
5	HENBA1005593	51.102	40. 350	125.688	37. 987	35.953	41. 577	39.834	47. 357
J	HEMBA1005595	52. 429	18.652	31.240	8.095	8,750	14. 586	12.433	7.264
	HEMBA1005597	125, 119	43, 335	90.414	24, 402	44.780	74.946	66.352	45. 322
	HEMBA1005606	141.646	66.667	95.041	30.084	57.974	107.329	84.655	46. 145
	HEMBA1005609	77, 991	60. 190	244. 951	52.002	41.602	40. 406	26.928	42.614
							32. 256		
	HEMBA1005616	47.760	62.865	190.870	42.670	41.809		23.683	43. 139
10	HEMBA1005621	33. 797	18, 993	22.515	11.333	11. 545	16.964	12.122	13.910
,,	HEMBA1005627	128.661	66. 487	148. 021	45. 359	42.161	42.054	30.884	43. 319
	HEMBA 1005628	43, 539	36.758	85.714	25. 524	46.601	19. 229	82.784	36.636
	HEMBA1005631	21.340	8.467	38.068	22.476	18.318	17.813	12.599	28. 199
	HEMBA1005632	113, 190	73.661	233.637	59.097	45. 388	52.090	29.944	37. 461
	HEMBA1005634	123.668	195. 912	390.579	101.523	107.528	72.729	54, 939	130. 473
	HEMBA1005662	15. 391	11.345	23.021	7.453	5. 561	13.084	8. 973	5. 282
15				34. 983	13.220	31.573	24.609	13.796	28. 043
	HEMBA1005666	33.844	30, 419						
	HEMBA 1005670	91.667	63.609	255. 523	57. 730	46. 927	45. 285	23.794	46.684
	HEMBA 1005671	63. 448	55. 388	34.948	26. 297	20. 567	2.367	5.666	13.509
	HEMBA 1005679	53.089	33. 284	126.705	39.666	32. 151	40. 446	37.522	36.817
	HEMBA1005680	115. 289	72.018	220. 408	76.653	55.707	68.735	32.613	36. 282
	HEMBA1005685	68.783	46.211	72.197	30.110	32.724	43.022	37.740	33.510
20	HEMBA1005698	37.890	35.679	44.793	29.794	25. 150	48.613	15.651	20.648
	HEMBA1005699	14. 243	17. 539	37.269	9.035	12.276	5. 454	5. 259	6.787
	HEMBA1005703	19. 524	15.116	20.249	7, 662	15. 489	11.648	8.488	10.229
				66.552	26. 492	29.605	90. 298	18. 303	44. 730
	HEMBA1005705	35, 316	35. 677						
	HEMBA1005712	20, 312	29.695	30. 267	17.829	17.668	17.695	14.517	23. 820
	HEMBA1005717	47.313	15.037	30.499	6.950	13.391	32.044	15.084	7.078
25	HEMBA1005718	88. 576	81.734	176.773	75.414	46.080	58.797	49.803	76.705
	HEMBA1005721	84. 981	42.340	58. 434	18.134	34. 246	43. 284	34. 523	41.460
	HENBA1005722	174.952	92.346	194.868	55.652	48.768	63.471	92.755	56.031
	HEMBA1005724	32.655	8. 284	5, 342	4.000	14.801	15.671	9. 324	5. 953
	HEMBA1005732	89.624	24. 907	32.546	5.638	21.753	30.046	28. 487	20. 595
	HEMBA1005737	25, 179	16.797	16.017	10.703	12.731	12.444	8. 579	7. 257
	HEMBA1005742	11.547	23. 162	24.345	20.921	29. 934	18.597	13.749	22. 702
30	HEMBA1005745			21. 907	16.923	13. 431	12. 235	10.908	8.606
		36.098	14. 407						
	HEMBA1005747	80.718	30. 396	44.843	21.861	30. 274	80. 588	47.082	26.037
	HEMBA1005749	35.749	31.758	64.769	22.766	28.853	26.733	31.698	30.753
	HEMBA1005755	34.680	39. 133	30.663	37.837	21.308	24. 392	15. 905	25. 470
	HEMBA 1005760	118. 125	41.490	33. 276	25.724	28. 933	45. 295	36. 173	31.205
	HEMBA1005765	94. 451	70.516	200.826	48.023	37. 340	35.414	31.098	40.041
35	HEMBA1005766	112.861	70.359	87.247	48. 958	51.073	52. 147	72.391	63.859
	HEMBA1005780	55. 961	34.713	89.816	28. 466	46. 254	28, 283	25. 156	29.122
	HEMBA1005795	18.800	38. 386	19.666	10.007	13.009	11.811	13.106	14.493
	HEMBA1005809	67. 301	66.510	87.390	53.061	43. 975	35.574	35. 334	57.818
	HEMBA1005813	52.911	84.881	160.064	38.752	43.727	30.799	23.426	57.177
	HEMBA1005815	30. 398	30. 434	43.366	19.911	16. 123	39.746	26.743	28. 548
40	HEMBA1005822				51.932	30. 845	20. 187	22.641	29, 114
40		40.948	47.746	65. 298					
	HEMBA1005829	114. 982	70. 536	272.004	48.816	36. 558	40.259	23.443	35. 824
	HEMBA1005833	59. 540	25.743	29. 266	15. 545	24.711	26.964	17.968	18.807
	HEMBA1005834	151.440	82.917	322.413	102.348	74.711	59. 590	35.082	70.415
	HEMBA1005844	66.624	11.865	96.556	95.719	56.133	75. 546	55. 974	122.840
	HEMBA1005852	71.743	77.830	72.218	53.009	85.623	78.593	90.291	87.310
45	HEMBA1005853	52, 809	83. 326	343.381	63.897	79. 208	48. 939	27.359	58. 458
70	HEMBA1005878	139, 991	109.928	447.600	93.748	65. 325	53.917	35. 383	63.446
	HEMBA1005883	5. 211	6.310		14.769	10.070	6.635	4.486	11.850
	HEMBA1005884	9. 136	10.768		9.504	7. 302	9, 142	4.561	12.287
	***************************************		12. 500			7. 370	7.346	1. 250	5. 470
	HEMBA1005891	8. 927							35.934
	HEMBA 1005894	70.006	59. 347			29. 584	23. 227	14.651	
50	HEMBA1005898	84. 399	61.254			43. 955	25, 491	23.019	41.130
	HEMBA1005902	38. 306	16.873			33.189	39.317	26.778	43.681
	HEMBA1005907	4.806	3. 997	8. 804	5. 339	3. 957	17.078	5.311	4. 941
	HEMBA1005909	4, 140	3.733	23.479	2.443	4.661	6.683	0.750	10.643
	HEMBA1005911	143.926	92.613			51.954	60. 593	39.302	55. 189
	HEMBA1005912	18.801	17. 269	13.568		21.976	14.454	12.917	26.318
	112-01:1000312	1 .3.001	1 503		1				

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Table 35

	(UEUG) (AAPA			4. 466 [36 655	- A 47A T	7 105 1	7 007 1	12 010
	HEMBA1005913	10. 533	16.117	14. 368	16.655	8. 179	7. 135	7. 907	12.918
	HEMBA 1005921	83. 262	45. 648	252.573	51.044	41.764	22. 286	23. 762	46. 202
_	HEMBA1005922	64. 440	17.427	35. 136	20.084	33.779	24. 835	18. 394	14.883
5	HEMBA1005929	173.002	139.696	378.444	96.543	83.075	72. 298	55. 205	94.716
	HEMBA1005931	146.354	89. 551	224, 601	83.623	63.406	73. 122	54. 973	59. 891
	HEMBA1005934	141.558	91.791	227.012	89.834	99. 341	96.876	62. 967	55. 492
	HEMBA1005945	144.693	21.871	18, 980	19.915	46.699	78. 590	80. 430	30.052
					21.004	17. 931	29. 331	21. 199	20.008
	HEMBA1005962	67. 209	34.719	63.745				7. 674	
10	HEMBA1005963	18.320	6.954	9. 127	5.913	2.497	8. 674		4. 873
,,	HEMBA1005990	581.646	117.336	139.967	53.671	242.262	424. 182	418. 873	85. 511
	HEMBA1005991	67.437	59. 327	188. 570	42.994	21.101	33.868	19, 164	11.619
	HEMBA1005999	193.878	135.695	450.789	126.399	129.150	103. 289	53. 193	115.911
	HEMBA1006002	73.560	26. 438	22.156	12.657	16.731	16. 116	10. 500	19. 305
	HEMBA1006005	59.620	7.083	16.863	8.213	29.019	53. 513	52. 130	23.838
	HEMBA1006011	25.811	30.413	39.888	21.434	54. 488	30. 978	27. 996	25. 339
15	HEMBA1006013	51.604	13.251	19.743	11.817	15. 364	28. 363	21. 493	18.674
	HEMBA1006016	101.929	42.149	115.996	36.228	39.875	46.607	33. 305	26. 397
	HEMBA1006019	31.772	18. 482	22.979	15. 207	22.984	24. 244	26, 246	14. 100
	HEMBA1006021	26. 984	10.213	45. 937	9. 253	20.615	14. 587	14. 203	12. 296
							46.057	25. 008	19. 323
	HEMBA1006022	100. 930	40.046	62.368	42.744	23.660			5. 133
	HEMBA1006031	42.088	41. 281	14. 729	11.264	12.725	36.716	13. 037	
20	HEMBA1006035	10.089	10.059	27. 290	8. 123	6. 309	6.629	2. 039	5. 229
	HEMBA 1006036	188. 431	82.469	443.914	119.939	80.135	81.126	54. 157	94. 631
	HEMBA1006042	69.906_	33.773	134. 462	30. 108	23. 244	29. 765	29, 479	29. 607
	HEMBA 1006044	53.721_	10.199	12.818	4.725	8. 467	5. 436	2. 586	4. 088
	HEMBA 1006045	48.078	43.730	61.128	28.336_	25. 311	26. 461	23. 478	44. 272
	HEMBA 1006048	35.685	18.435	41. 495	19. 225	19.636	34. 213	26. 302	28. 809
25	HEMBA1006053	0.000	356.500	78.844	24. 270	47.030	114. 986	63. 574	385. 970
25	HEMBA1006055	7.603	5. 331	12.625	4.484	13.776	12. 227	9. 079	5. 545
	HEMBA1006058	51.872	19.394	14.828	7.834	11.877	25. 640	15, 830	21.486
	HEMBA1006063	72.886	52.429	63.882	34.021	30, 125	39. 536	28. 303	35. 860
	HEMBA1006067	6.005	14. 253	7.505	3.169	2.242	3. 352	4. 358	0.888
	HEM8A1006081	70. 282	19.151	25.838	8.981	9. 908	26.560	16.837	23, 976
	HEMBA1006089	54. 392	23.145	42.709	18. 278	17.433	17.768	18. 372	23. 981
30	HEMBA1006090	71.092	20.389	36.832	15. 386	17.868	38.904	35, 031	18. 238
	HEMBA1006091	69.022	28. 947	126.425	16. 353	30.302	56.034	53.660	66. 468
		111.885	11.435	50.738	16. 185	27.687	43. 178	26.048	14. 980
	HEMBA1006093		27.136	19.149	18. 199	31.100	31. 158	28. 536	25. 484
	HEMBA1006099	40. 381						13. 379	34. 466
	HEMBA1006100	36.979	48. 991	259. 267	41.090	50.094	24.833		
0.5	HEMBA1006108	40.170	19.301	21.811	11.126	8.795	12.441	8.780	16. 453
35	HEMBA1006114	42.849	44.783	46.702	33. 193	23. 220	34.626	28. 294	51.756
	HEMBA1006121	160.208	21.943	26.728	10.160	21. 331	17.129	26.838	25. 137
	HEMBA1006124	63.151	11.764	15.994	17.764	14.099	57. 249	29. 200	8. 240
	HEMBA 1006 125	72.730	70.406	57.020	50.057	45. 287	40.856	45. 665	68. 939
	HEMBA 1006 130	36.221	31.688	34.742	7.817	28, 246	34. 473	25.726	21.315
	HEMBA 1006 138	160. 258	170.815	435. 120	106.719	139.660	100. 947	67, 854	89.604
40	HEMBA1006142	127. 194	85. 725	238. 562	54. 531	52.936	65. 032	45. 938	59. 791
	HEMBA1006150	66.777	58.231	75.666	59. 941	19.605	46. 114	33. 261	75. 731
	HEMBA 1006 151	189. 265	57. 959	104. 921	29. 646	46.546	66.736	74. 155	88. 383
	HEMBA1006155	141.288	19.560	50.142	11.752	32.711	79.435	60.621	32.838
	HEMBA1006158	17.276	12.039	19, 210	7. 139	7.468	23.241	7.360	13. 357
	HEMBA1006164	140.272	70.843	382.965	97.488	87.832	69.460	42. 210	85. 135
	HEMBA1006171	66.839	48.304	34,618	13.911	21.700	40.783	26.049	37. 233
45	HEMBA1006173	63.939	35. 393	52. 598	22.894	32.403	35, 413	40.872	67.870
	HEMBA1006176	51.671	222, 661	52.703	39. 369	29.305	59. 271	24. 272	83. 343
	HEMBA1006182	72.842	38. 362	132.455	29.730	26.735	30. 382	19, 907	34. 405
				37. 528	55. 808	23, 143	18.848	13. 456	40.765
•	HEMBA1006197	16.655	31. 338					14. 389	
	HEMBA1006198	30.466	15. 178	21 337	16. 185	25.764	15.643		18.561
50	HEMBA1006213	38.783	20.120	38. 136	15. 627	10.604	25.761	21.716	35. 282
•••	HEMBA1006217	12.003	18.510	33,960	4.079	17.107	31.016	36. 526	19.419
	HEMBA1006226	40.304	60.090	110.529	40.359	39.915	62.796	35, 202	59. 281
	HEMBA1006235	40.954	9.021	21.361	7. 280	14.241	13.056	4.951	7.077
	HEMBA1006248	42.946	17. 521	32.092	10.747	12.992	19. 331	18. 339	17. 999
	HEMBA1006251	84. 944	24. 303	30.554	15. 291	24.212	30.870	18.154	10. 996
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Table 36

				140)IC 30				
	HEMBA1006252	36.069	24.612	74, 170	29.506	28, 055	19.517	14.085	15. 356
	HEMBA1006253	75.854	7.002	20.773	16. 455	11.705	12.936	6,506	11.398
	HEMBA1006259	37.456	48, 402	136,000	39.735	19.462	25. 242	19.832	25. 931
5	HEMBA1006261	23.677	23. 578	6.874	13.012	7, 127	69.427	7.141	17. 143
	HEMBA1006268	35.886	12.563	30.879	8.970	7.077	19.793	22.288	18. 289
	HEMBA1006271	122.980	98.618	185.469	77.610	45. 268	47.910	36.533	47.867
	HEMBA1006272	16. 261	12,829	9, 416	4. 968	5. 925	27,766	15.997	7. 567
	HEMBA1006273	47.890	12.641	71.219	20, 880	30.446	30.473	18. 419	22. 459
	HEMBA1006276	79.296	11.878	30.854	34. 032	8.760	27. 168	16. 165	7. 501
10	HEMBA1006278	40.093	7.717	26.091	4. 506	18.669	11.680	11.224	9.893
	HEMBA1006283	16, 994	23. 586	25.614	25. 226	23. 447	59.086	28. 267	25. 848
	HEMBA1006284	29. 982	22, 166	27. 891	20.874	8. 594	18. 386	15. 293	13. 396
	HEMBA1006291	22.745	13.071	36.861	9.670	4. 059	11.649	31.851	7. 519
	HEMBA1006292	17, 718	8.916	20.081	10.169	4, 378	7. 903	9. 259	7.898
	HEMBA 1006293	31.307	10.056	8.749	4,645	4.097	6.631	8. 473	7. 189
15	HEMBA1006299	21, 091	5, 917	6. 157	1, 371	4. 543	2.465	1.701	2.648
	HEMBA1006309	69, 975	25. 568	110.869	33, 191	19.510	31.160	24.850	17.764
	HEMBA1006310	40.983	23. 265	36. 585	20.570	11.748	29.056	27. 263	17. 748
	HEMBA1006311	85. 398	20.844	64.711	8. 925	20. 171	92.798	9. 481	19. 313
	HEMBA1006313	27.762	12.975	47.707	17, 417	7. 455	13.117	9. 891	6.082
	HEMBA1006316	23.345	3.751	3.303	2.158	8.774	9.668	8. 505	3. 270
20	HEMBA1006328	79.937	83.744	185. 981	41, 111	28.820	37.527	35. 377	85.968
20	HEMBA1006334	22.524	16.717	17.679	5.994	8. 506	9.813	3.866	- 5. 361
	HEMBA1006335	72.666	41.477	35. 235	27.435	6.110	5.851	24. 375	8.434
	HEMBA1006344	34.707	67.866	132.978	46.518	34.812	40.158	41.934	25. 330
	HEMBA1006347	34, 301	16.445	32.190	19.603	16.749	20.762	20.884	15, 376
	HEMBA1006349	139.389	26.300	48.767	43. 275	22.026	24.648	22.876	21.499
0.5	HEMBA1006352	21. 127	17.873	15. 526	9,410	8.472	14.845	7.491	9.414
25	HEMBA1006357	94. 337	82.319	287. 531	67.888	76. 120	47. 179	41.500	59. 557
	HEMBA1006358	48. 925	31.345	132.494	32.473	25.019	28. 197	13. 250	24. 899
	HEMBA 1006359	57. 203	18. 522	160.314	70.923	17.441	30.686	11.154	47. 991
	HEMBA1006360	29.518	10.133	15. 515	17. 275	6. 141	13.876	6.804	8.361
	HEMBA1006364	59. 236	7. 900	27.522	12.114	5. 401	15. 432	17. 981	6.672
^^	HEM8A1006377	67.120	31.113	57. 269	33. 567	23.849	45. 246	31.609	20. 280
30	HEMBA 1006380	73.227	57.029	182. 581	57.870	22. 288	33.416	23.616	40. 932
	HEMBA 1006381	359. 346	122.755	376.090	126.304	112.826	145.346	91.469	93. 252
	HEMBA 1006385	60.234	62.166	257. 945	59. 429	59.157	40.136	35. 385	17. 281
	HEMBA 1006390	71.393	38.752	46.828	25.848	16. 455	41.253	16.013	27.609
	HEMBA1006391	61.261	18.765	20.686	10.972	10.022	39. 431	27. 305	11.797
0.5	HEMBA1006398	42.089	3. 225	18.036	5. 299	25. 386	6.480	0.000	3. 308
35	HENBA1006405	137.413	28.645	40. 904	17.896	18. 180	84. 926	41.325	24.773
	HEMBA1006410	149. 580	32.840	61.022	20.027	39.718	54, 551	23.826 37.809	33. 928 33. 447
	HEMBA1006416	96.031	62. 892	198.896	50.538 11.378	38. 551 10. 280	37. 025 28. 208	46.245	36. 223
	HEMBA1006418 HEMBA1006419	23. 236 189. 293	18. 335	476.145	90.626	79. 213	64. 306	40.042	52. 384
	HEMBA1006421	39, 702	26. 487	127. 221	23.773	16. 184	14. 460	12. 270	13. 523
40	HEMBA1006424	4. 484	36. 452	10. 588	3.778	4. 512	7. 346	2. 324	3. 323
40	HEMBA1006426	88.597	67. 224	230.530	60.836	32.273	40. 489	17, 284	36. 244
	HEMBA1006430	61.672	17. 989	69, 151	15, 913	11.038	15.595	9.696	17.632
	HEMBA1006438	45.084	34. 475	111.512	27.012	15.035	34.111	12.678	11,056
	HEMBA1005445	48. 245		53.981	9. 326	15.672	34.167	27.442	18.331
	HEMBA1006446	22.911	3.160	3. 324	1.568	4. 341	2.585	1.331	0.000
45	HEMBA1006456	36.915	28.165	141.114	18.927	65.823	33.549	13.651	33. 405
45	HEMBA1006461	60, 747	42.392	161.108	40. 447	22.274	32.823	18.018	27. 165
	HEMBA1005467	13, 357	6.130	15.734	10.759	4.032	4. 471	6.183	2.655
	HEMBA1006470	73.960	30.706	103. 625	27. 235	29.870	33.756	33.818	24. 286
	HEMBA1006471	19.032	4.504	7.503	2.933	2. 522	5. 224	10.020	1.873
	HEMBA1006474	25.718	12.420	21.381	11.498	9.614	19.875	17.655	13.491
	HEMBA1006476	180.042	91. 936	63.588	43. 462	42.248	109.725	88.725	65. 945
50	HEMBA1006482	129.627	169. 312	167. 982	151.338	57.839	95. 521	75. 480	239. 325
	HEMBA1006483	99. 620	64.773	232.207	50. 445	29.074	37. 572	23.818	27.130
	HEMBA1006485	41.690	4. 055	17.445	11.682	4. 522	9. 351	6.411	10.066
	HEMBA1006486	76. 250	36, 421	29, 634		17. 302	21.229	17.832	15.706
	HEMBA1006489	5.771	32.673	2, 141	5. 240	2. 356	4. 324	4.739	7. 328

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Table 37

									10 000
	HEMBA1006492	14.002	19, 916	24.662	35. 451	8.836	8. 075	11.419	12.090
	HEMBA1006494	7. 279	0.000	19. 790	3.750	8.718	8. 343	5.851	5. 887
_	HEMBA1006497	41.284	12.396	23. 326	6.590	7.186	11. 228	9.062	5.781
5	HEMBA1006501	160.565	16.895	26.893	13.446	17.608	65. 467	41.560	6. 197
	HEMBA1006502	53.451	19, 114	39, 593	25. 366	10.919	15.054	17.536	15.658
		19. 274	8. 180	10. 287	4. 521	7.939	5. 288	15. 480	10.062
	HEMBA1006507				18.732		45. 881	29.819	
	HEMBA1006517	95.989	30.085	91.871		21.918			16.672
	HEMBA1006521	31.224	27.873	37.864	18.318	9.774	14. 205	14.646	13.907
40	HEMBA1006529	28. 702	20.010	34.050	20. 150	16.588	7.353	8.993	17. 327
10	HEMBA1006530	18.445	16.411	29. 175	14. 433	12.214	15.734	15.731	8.081
	HEMBA1006535	11.627	7.208	18.048	3.956	8.160	19.824	5.837	3. 457
	HEMBA1006536	68.087	40.009	142.475	43.263	34.343	42.050	42.157	23.975
	HEMBA1006540	20. 393	10.867	35, 153	8.637	8.656	15.027	11.094	10.350
	HEMBA1006544	30. 281	4, 662	59. 940	7. 791	7. 169	15.883	8.745	8.693
	HEMBA1006546		53. 155	127. 193	49. 337	73.807	60.506	22.328	34.045
15		68.722					14, 211	8. 987	6. 080
10	HEMBA1006549	13.885	13.666	21.800	11.666	8. 491			
	HEMBA1006559	26. 976	22.040	38. 197	16.910	14. 550	14.058	13.018	17. 217
	HEMBA1006562	55. 924	24.663	75.789	20. 363	17. 181	26.651	18.158	19, 510
	HEMBA1006566	20.849	6.116	14. 933	8.767_	9. 572	6.937	5. 229	4.788
	HEMBA1006569	57.508	20. 299	44. 291	27.048	12.798	15. 243	24.739	31.861
	HEMBA1006572	21.817	4. 339	15.862	1.796	3.407	11.582	8.381	5, 922
20	HEMBA1006579	5. 427	18. 336	4.219	3.440	2.139	5. 460	3.967	5.110
20	HEMBA1006583	31.967	15.854	29. 307	14. 271	11.747	26.889	17.058	10.451
	HEMBA1006595	59.014	41. 577	148. 359	30.660	16.681	19.571	13. 265	24.768
				210.001	47.574	27.392	47.009	27.887	28.666
	HEMBA1006597	111.817	64. 480		41.314				
	HEMBA1006606	79. 184	47.311	131, 822	40.177	33. 228	35.403	25. 240	31.687
	HEMBA1006612	43. 105	20.909	46.913	39. 205	20. 348	25. 383	18.706	17, 150
25	HEMBA1006617	79.139	62.924	235. 236	60.258	30.407	40.264	28. 184	38.643
	HEMBA1006624	449. 384	84.050	165.494	39, 352	209. 908	291, 427	208. 533	65.478
	HEMBA1006631	168. 309	108.316	381.778	89.696	71.812	80.634	39. 325	50.996
	HEMBA1006635	51.406	33.730	158. 286	28.605	19, 347	19.781	9.639	12.894
	HEMBA1006639	67.363	30.354	51.867	15.409	33.210	43.083	25. 295	12.985
	HEMBA1006643	229.685	30.246	56.218	16, 406	35, 196	68.642	41.724	17. 931
	HENBA1006648	80.985	32.464	39, 607	14. 926	36.718	12.135	32.217	48.853
30	HEMBA1006652	118.455	69. 232	231, 917	50.609	51.023	50.716	21.698	29.527
	HEMBA1006653	46. 971	16.614	46.472	16.579	12.358	15.364	13.867	9.224
		89.823	28. 363	60. 976	37.660	28. 124	47.014	33.470	16.872
	HEMBA1006658				49. 132	29. 124	33.070	25. 182	33.784
	HEMBA1006659	79.863	33.626	48. 217				12.046	10.419
	HEMBA1008665	25.726	26.740	39.661	13.975	13. 287	15. 240		
	HEMBA1006665	8, 276	4. 281	10.565	6.319	4. 257	10.392	2.791	2.171
35	HEMBA1006671	39. 553	178, 623	135.413	18, 941	17. 294	37.782	10.166	32.048
	HEMBA 1 006674	100.472	44. 108	176.724	46. 922	36.367	44.809	43.576	43. 269
	HEMBA1005676	120.417	42.888	163.816	29. 504	40. 435	60.162	32.540	34. 825
	HEMBA1006682	27. 104	2.556	23. 174	4. 035	8. 982	19.092	3.958	0.000
	HEMBA1006688	57. 351	56.288	111, 358	60.597	65. 322	37.545	20.757	20.789
	HEMBA1006695	132.496	140.334	315.655	97.296	56.206	54.392	37.622	57. 596
40	HEMBA1006696	65, 136	25. 204	42, 137	26.654	26.490	30.156	6.159	27.512
40	HEMBA1006702	4. 275	4. 328	8.881	7.114	3.362	1.846	7.796	1.965
	HEMBA1006707	52.417	20.766	26.862	21.409	19.843	32.229	13.146	18.546
	HEMBA1006708	126.875	38. 520	66.803	31.253	33. 294	55. 347	32.071	18. 229
				94, 432	24. 924	17.365	30. 329	18.603	23.474
	HEMBA1006709	67. 500	31.686		12 664		54. 326	26.752	11.544
	HEMBA1008717	110.641	21.536	29. 255	12.664	16.091		10 505	
45	HEMBA1006724	34. 421	23.073	25.607	18. 231	12.305	27.570	10.585	17.797
	HEMBA1006731	36.072	18. 255	41. 441	15. 382	16.479	17.272	10.826	15. 482
	HEMBA1006737	50.467	14. 107	30.096	14. 542	20. 232	22.606	10.316	11.440
	HEMBA1006742	60. 258	45.190	134.964	35. 452	21, 315	21.889	15. 223	23. 529
	HEMBA1006743	41.970	22.864	31.760	22.024	15. 126	23.989	13.179	16. 281
	HEMBA1006744	181.068	97.273	433.004		69.785	59.354	46.770	61.806
	HEMBA1006749	51.776	9.753	37, 994		23, 164	34.516	28.426	23.238
50	HEMBA1006752		50.318	88, 111	59.765	47. 490	69, 461	37.541	47.074
		124, 800				17.745	16. 269	10.783	12.424
	HEMBA1006754	49. 957	30.459	85.726	23. 747				15.933
	HEMBA1006758	75. 460	21.737	26.190		18. 249	38. 492	30.654	
	HEMBA1008767	14.002	15. 106	11.961		5, 628	13. 334	8. 382	8. 573
	HEMBA1006770	120.485	21.505	62.144	29.559	32.512	49.739	45. 952	28.318

Table 38

,	UE POLICE TO			100 029	47 164 1	20 155	26 722 1	10 025	29, 256
	HEMBA1006779	81. 492	51.077	162.657	41.163	39, 166	36.722	18. 025 41. 524	39. 494
	HEMBA1006780	78.359	78.052	345. 442	73.371	68.858	55. 888	19.736	25. 701
	HEMBA1006789	29. 455	21.233	20.440	14. 349	11.547	38. 549		40.750
	HEMBA1006795	143.727	88. 701	218.732	55.068	49.500	46. 284	21. 141	
	HEMBA1006796	87. 214	15.814	115. 542	17.685	16.790	38. 694	15. 525	15. 352
	HEMBA1006805	68, 116	31. 212	153.041	33. 162	30.301	34. 197	24. 275	30.733
	HEMBA1006807	94. 524	86.723	157.559	64. 349	36. 505	62. 933	23.097	55. 508
	HEMBA1006813	40.696	4. 415	4.750	4. 264	10. 978	7.562	6, 201 20, 346	3. 198
	HEMBA1006819	53.717	15, 217	30.071	14.679	17.006	30.866		8. 250
	HEMBA1006821	39.052	30. 425	111. 325	35.769	34. 975	22. 216 27. 255	18. 924 21. 689	20. 698 27. 074
	HEMBA1006824	68. 491	61.498	201.721	47. 107	40. 322 40. 147	75. 996	66.799	71,706
	HEMBA1006832	84.462	89. 500	102.038	77.046 48.909	41.460	61.443	30. 402	31. 940
	HEMBA1006834	123. 958	57. 085	160. 407 38. 470	23. 193	18. 979	22.344	22, 426	16. 742
	HEMBA1006835	33. 705 52. 436	19. 529 44. 642		258.615	195.878	33. 141	8.256	13. 117
	HEMBA1005843		34. 224	158. 388	39.483	30.349	34, 943	15.743	28. 240
	HEMBA1006849	88. 931 44. 733	24, 923	67.667	24. 186	15. 829	36.593	11.223	18. 454
	HEMBA1006850	215, 207	94. 180	158. 997	67.349	259. 512	135. 856	371. 932	44. 053
	HEMBA1006865	124. 996	59.773	124. 376	43.328	69.356	71.072	66.350	45. 129
	HEMBA1006867	15.632	11.094	39.646	14. 084	12. 902	11.855	5. 865	18. 338
	HEMBA1006873	9.965	9. 279	7.010	5.013	6. 262	5. 127	7. 141	8. 422
	HEMBA1006877	44. 043	18. 321	20. 546	8. 172	14.670	13. 165	16.493	9, 073
	HEMBA1006878	100. 427	34. 418	109.029	25. 739	29. 525	48.800	41.513	17. 905
	HEMBA1006879	108. 299	42.811	121.051	60.872	47.507	40.075	14. 429	51.924
	HEMBA1006884	95. 426	29. 331	67.556	27.787	25.909	106.818	47.878	47.793
	HEMBA1006885	107.720	54. 342	127.920	62.272	55.739	51.739	36.790	50.612
	HEMBA1006886	50. 841	22.970	51.528	12.561	20.660	23. 207	26. 952	19.149
	HEMBA1006889	81.809	20. 952	21.474	12.691	24.681	41.822	48.768	15. 196
	HEMBA1006896	68. 030	97. 285	75.370	52.746	23.109	44. 481	37.701	50.662
	HEMBA1006900	61.515	36, 410	61.016	23.329	21.390	38. 404	27. 583	22.774
	HEMBA1006902	43. 283	19. 713	47. 129	12.105	11.602	27.830	26.548	13.885
	HEMBA1006912	183.904	90. 995	338. 160	78. 230	79.588	63.729	39.994	64. 953
	HEMBA1006914	54. 548	39, 053	48. 945	35.736	25. 895	38. 585	22.479	33.810
	HEMBA1006916	62. 872	0.000	65.115	29.982	32. 625	61.537	62.750	30.818
	HEMBA1006921	64. 867	21.840	74.902	15.692	30.866	41. 257	25. 569	10.362
	HEMBA1006926	51. 195	10.616	76.671	24. 435	20.300	84. 402	29.503	20.967 7.086
	HEMBA1006927	24.016	13.778	23.573	5. 335	15.250	11.291 5.970	11.672 5.688	3, 134
	HEMBA1006929	7. 146	8. 487	5. 431 45. 566	5. 526 20. 391	16.346	25. 493	20, 196	17.720
	HEMBA1006936	68. 233	22. 847 8. 409	31.234	7.743	5. 002	6.780	6,773	5. 945
	HEMBA1006938 HEMBA1006941	14. 202 30. 559	24. 290	40. 928	13.779	16.040	34. 253	22.542	18.507
	HEMBA1006942	147. 487	57.842	121.883	69, 207	55. 456	76.853	61.942	66.640
	HEMBA1006942	80. 546	64. 930	104.037	63.709	40. 444	54.676	33.533	31, 915
	HEMBA1006949	10. 292	41. 467	23. 921	1.860	15. 813	7. 071	10.866	5. 231
	HEMBA 1006952	58. 685	12.572	34.750	8,032	18. 283	39.764	15. 332	12, 456
	HEMBA1006960	91. 939	38. 895	93.164	24.834	34. 400	36. 160	36.715	34.791
	HEMBA1006973	74. 208	24. 793	50.621	17.619	22.844	24. 971	24.844	16.167
	HEMBA1006974	48. 691	39.013	59.414	48.064	16.799	38. 579	21.301	46.006
	HEMBA1006976	35. 907	15.675	32.116	19.091	14. 522	30.574	25. 042	18.348
	HEMBA1005989	6. 422	2. 207	2.374		2.670	3.696	2.557	3. 536
	HEMBA1006993	334. 266	64. 150	357. 947	46.138	95. 466	144.777	109.174	54.000
	HEMBA1006996	9. 183	9.870	15.032			9.518	8.368	9.637
	HEMBA1007001	117.610	95, 668	334.868		55. 288	47.863	27. 205	56.828
	HEMBA1007002	93. 134	41.846	72.311		16. 249	59. 722	46.434	40. 628
	HEMBA1007013	65. 73.4	23. 106	53, 712		20.783	34. 293	29.163	29. 338
	HEMBA1007016	36. 649	14. 972			9. 597	17. 982	16.658	15.035
	HEMBA1007017	6, 290	0.000				2.329	1.949	0.000
	HEMBA1007018	19, 457	15.664			10. 586	15.084	9. 105	14. 124
	HEMBA1007044	139.784	50.078			53.729	123. 367	90.838	36.173
	HEMBA1007045	49. 576	7.913				19.099	12.683	7.276
	HEMBA1007051	36. 374	44. 117			19.407	24. 088	15.546	9.363
	HEMBA1007052	69. 582	19.611	40. 507			19.409	18.969	10. 939
	HEMBA1007053	25, 326	27.611				20.128	7.847	9.544
	HEMBA1007057	45. 897	13, 545	33.857	18.616	25. 861	36.241	14.769	13.902

Table 39

	TIELEDA LOGRADA	1 100 010							
	HEMBA1007062	129.012	18.903	40.670	21.323	29.469	40. 252	29.408	15.700
	HEMBA1007063	81.681	45.884	187.380	52. 391	35.943	28.608	35.303	41.236
	HEMBA1007066	98.395	32,970	35. 373	22.961	11.085			
5							42.430	26,631	14.760
•	HEMBA1007069	23.449	21.519	78.409	16.835	27. 425	17.217	9. 095	16.163
	HEMBA1007073	54.833	42.548	40.682	29.352	11.879	7.937	24. 282	19, 372
	HEMBA1007076	83. 020	48.746	248.260	61.189	50, 193			
	HEMBA1007078						68.045	43. 836	35. 650
		151, 561	159.600	446.445	189.146	130.283	98.734	65.934	117.079
	HEMBA1007080	43, 963	44.765	174. 545	66.950	45.879	43.194	43. 909	50, 100
	HEMBA1007084	78. 948	60.672	268. 327	63.769	63.088	60.307	35.006	
10	HEMBA1007085								46.866
		263, 538	108.018	162.599	48. 155	77. 545	161.321	63.614	80.640
	HEMBA1007087	85. 598	25.085	47.862	25. 580	13.918	62.815	143.461	30.856
	HEMBA1007089	21, 131	32.023	21.145	14.738	7.213	19.681	9.036	10.026
	HEMBA1007095	147, 777	215, 051	136.910	63.992				
						170.706	117.992	103.152	86.452
	HEMBA1007101	78. 959	53.790	147.891	35.676	28.082	27. 200	19.131	25. 922
	HEMBA1007104	66. 308	23, 279	45.417	11.902	19, 468	48.054	26.760	16.647
15	HEMBA1007106	28, 449	17.761	41.268	28.670	17.681	14.174	10.999	
	HEMBA1007112	12.759							7.534
			8.412	16.340	9.319	7.661	7.304	13. 296	6.622
	HEMBA1007113	126. 702	0.000	229.408	64.551	40. 242	39.032	13.319	26.174
	HEMBA1007121	219.036	207.410	696.658	149. 217	168.827	131.628	642.099	128.755
	HEMBA1007129	50.726	42.510	63.847	31.663	26. 417			
							24. 371	18. 928	20. 103
20	HEMBA1007147	111.299	117.722	312.811	79.949	67. 395	74. 391	35.758	54. 184
20	HEMBA1007149	83. 453	6.442	19.831	7.332	11.043	9.349	9.831	8.756
	HEMBA1007151	97, 211	33, 530	53. 944	24. 544	18, 501	35. 246	36.228	24. 174
	HEMBA1007172	52.683	25. 324	438.704					
					42, 182	28. 599	38. 126	26. 167	25.770
	HEMBA1007174	52. 921	13.482	44.770	21.384	19. 520	28. 559	22. 332	20. 471
	HEMBA1007176	89.919	24.758	53.414	32, 841	44.643	73.679	87.040	30.762
	HEMBA1007178	93. 941	73, 120	135. 427	34, 313	32.040	34.622	22.898	24.897
25	HEMBA1007185	62.558	18.807	36,824				22.030	
					15. 490	20. 528	37. 568	22.260	12.783
	HEMBA1007186	70.967	31.546	59, 038	21.059	21.332	35.648	42.864	11.346
	HEMBA1007194	53.376	38.911	126.660	33.992	23.875	21.109	12.122	23. 307
	HEMBA1007200	74. 955	53.829	44, 212	23.979	20. 225	32.762	55.417	22. 176
	HEMBA1007203	87.803	26.807	41.357					
					14. 648	9. 791	23. 392	30.167	17. 274
20	HEMBA1007206	82,800	73.675	225. 293	44. 461	28.674	37. 091	14.673	34. 505
30	HEMBA1007224	25. 614	40.402	50.116	21.484	14. 920	22. 548	13. 197	20.053
	HEMBA1007226	88.512	43.606	93, 121	22. 209	17. 911	38. 704	43.759	31.721
	HEMBA1007240	131.657	62.804	86.650	9.510	21.890	53.116		
	HEMBA1007241	13 335						42. 250	16.655
		12. 225	7.719	18.461	5. 051	6.724	15. 945	3. 135	5. 390
	HEMBA1007242	21.409	14.030	13.648	11.068	6. 265	17. 370	8, 487	5. 236
	HEMBA1007243	61,824	25.854	40.264	17. 235	23. 438	39. 197	31.904	20. 347
35	HEMBA1007251	37.660	16.946	37, 149	16.699	12. 180		30.321	
	HEMBA1007256	53, 905					19. 482		10. 262
			43.642	113, 110	31.642	27. 946	30. 492	18. 548	23. 645
	HEMBA1007267	80.741	40.085	207, 160	61.174	38. 220	29.008	32.292	29.672
	HEMBA1007273	41.062	9.087	11.906	5. 193	6. 445	7.723	9. 225	4. 483
	HEMBA1007279	54, 376	20.734	133, 494	27. 987	21.355	19.941	17.364	19.503
	HEMBA1007281	8. 523	5.717	4,731					
40	HEMBA1007283	25, 940			3. 403	2.317	2.497	2.740	0.000
40			14. 444	24. 974	23.487	19.771	23.418	19.378	26. 409
	HEMBA1007288	57. 959	39. 576	155, 227	28.725	24.689	25. 110	16.998	16.095
	HEMBA1007291	37. 974	19.069	59. 253	20. 445	13.404	17. 376	13.060	13. 147
	HEMBA1007299	446.640	93.668	199.852	61.423	94. 129	249.345	241.373	85. 323
	HEMBA1007300	103.752	25. 694						
				24, 914	18. 217	40.413	26.018	31.407	16. 569
	HEMBA1007301	49, 752	18. 178	32.677	18. 170	33.650	33.786	22.892	12. 782
45	HEMBA1007319	13. 312	10.598	23. 453	16.511	4. 278	9. 382	2.996	8.570
	HEMBA1007320	53.723	23, 595	62.301	29.439	16.672	32.932	28. 191	18.418
	HEMBA1007322	45. 986	125, 362						
	DCDDLIGGTAGA			77.545	43.693	17. 955	45.689	39.556	80.836
	HEMBA1007323	64. 720	16.869	22.970	11.238	11.687	32.209	25. 350	7. 506
	HEMBA1007326	313.094	189, 188	862.276	214.045	178. 109	171.587	70.819	115, 174
	HEMBA1007327	78.767	61, 102	219.980	55.002	29.411	44.095	29.354	42. 286
50	HEMBA1007332	71.516	9.318	34, 879					
50					5. 559	7.452	24.826	12.763	20.050
	HEMBA1007341	89.805	53.431	207. 395	82.402	105.877	47.861	32.826	50.162
	HEMBA1007342	22.063	17. 289	28. 253	18.196	17.751	26.378	13.820	9. 173
	HEMBA1007347	112, 392	64. 499	230.022	60.348	47. 557	63.758	30.683	
	HEMBA1007353	1.685							33. 285
			3. 520	0. 575	1.860	1.976	0.107	2.061	0.788
	HEMB81000005	60.047	46.027	121.870	38. 241	20.699	18.268	20.068	26.957
55	HEM881000008	97. 929	53,604	274. 179	68.681	38. 935	39. 328	26.881	34.873
									37.013

Table 40

	HEMBB1000018	122.130	127.861	329. 165	120.419	57.867	95. 203	75. 902	92. 924
	HEMB81000024	181.606	97.019	373.954	102. 401	70. 406	70, 591	40.304	66.798
5	HEMB81000025	85. 919	29.049	45.055	23.789	13. 946	24. 397	29.349	13.072
v	HEMBB1000030	108.167	68.316	303.677	83.010	68. 378	81.687	34, 886	37.617
	HEMB81000036	107.960	11.573	50, 484	11. 277	20. 480	41. 381	25. 378	14.730
	HEMB81000037	77.688	29, 380	69.658	56. 679	27.020	54.062	30.086	15. 311
	HEMBB1000039	52.550	48, 503	140.795	30.096	18. 739	26.012	15. 151	21.723
	HEMBB1000044	134.136	75. 469	218.667	61.596	32.667	29.659	43.360	42.831
10	HEMBB1000048 HEMBB1000050	17. 937 74. 210	21.052	31.004 207.484	18. 291 35. 691	11. 321 22. 905	20. 120 25. 584	21.506 18.572	15.078
	HEMB81000054	68. 273	33.681 47.191	246.350	44.008	24. 522	29. 259	22.570	17. 494 21. 316
	HEMB81000055	72.875	112. 284	61. 172	110.297	21. 358	70.636	93.824	132. 288
	HEMBB1000059	331.577	184.687	662.540	182.481	130.065	131.364	90.002	121. 903
	HEMBB1000072	240.733	98.890	326.893	75.919	67.742	118.222	108.108	91. 458
15	HEMBB1000081	23.738	27.174	85, 100	21.146	30.856	20. 458	7.513	15. 351
75	HEMB81000083	120.759	58. 163	188. 224	40.609	37.789	59. 334	33.712	39. 101
	HEMBB1000089	67.618	54. 952	191.832	56.629	24.609	36.847	30.680	26.912
	HEMBB1000094	355, 534	116.828	161.958	31.504	29. 300	49.613	36.239	35. 197
	HEMBB1000097	27.834	63.724	51.488	14. 249	22.834	34, 068	18.547	16. 455
	HEMBB1000099	157.641	91.912	456.470	71.078	50.739	64. 471	32.108	43. 354
20	HEMBB1000103	75.781 62.814	59. 392 44. 996	114. 974 77. 918	44. 216 35. 044	31.915 19.825	47.628 40.409	23.669 26.156	56. 268 45. 001
	HEMBB1000113	43.660	33, 435	95. 987	42.744	19.714	20, 114	15.899	21.606
	HEMBB1000119	57.350	21.211	42. 528	17.770	19.517	28.754	23.570	30, 104
	HEMBB1000133	92.950	65. 230	58.619	69. 544	53.706	104. 229	39.058	80, 858
	HEMBB1000134	44, 120	20.654	76.693	40.611	24.712	37.185	42.327	21.963
	HEMB81000136	21.810	7. 191	44.517	15. 599	7.339	22.582	12.399	24.899
25	HEMBB1000141	163.867	99. 946	331.822	95. 807	55.858	64.560	36.737	52.602
	HEMBB1000144	96.831	97.019	183.423	88. 529	36. 185	15, 577	29.259	32. 144
	HEMBB1000147	59.253	9.088	62.426	7. 391	11.451	7.175	11.502	10.693
	HEMBB1000152 HEMBB1000154	56.391 85.308	28.723	34. 597 101. 061	15. 309 33. 881	19.424	32.469 27.298	29.105 20.174	19.117 15.366
	HEMBB1000154	35.691	47.878 36.132	109.038	28. 164	29.608	22.283	16.557	17.041
30	HEMBB1000173	170.611	173.001	494. 253	143.666	83.705	123.932	65.317	76. 388
	HEMBB1000175	32.273	19.114	23.481	10.948	4.039	29. 180	7.135	13. 322
	HEMBB1000176	56.984	51.334	90.749	69.004	40.144	52.980	25.845	19.359
	HEMBB1000198	70.426	12.768	26.381	10.237	6.266	11.215	8.858	5. 363
	HEMBB1000208	42.474	8.966	34. 929	10.418	12.883	9, 285	12.335	7.978
25	HEMB81000209	43.846	10.700	9.943	10.934	8.858	12.135	9.049	4. 168
35	HEMBB1000212	27.532	12.579	76.077	15. 361	33.518	17.471	13.132	16.552
	HEM881000215 HEM881000217	178. 324	89.053 45.416	294.606 96.614	95. 420 47. 569	68.598 37.572	89.720 89.989	51.270 48.073	61.235 33.510
	HEMBB1000218	88.298	123.000	347.859	84. 124	41.828	57.417	21.147	34. 605
	HEMB81000226	70.693	14.949	41.586	31.786	30. 261	28. 577	14.779	27.177
	HEMBB1000230	28.681	8.910	13.549	5. 500	3.547	9.616	6.632	3. 293
40	HEMBB1000240	44.662	12.588	13.211	10.455	4. 589	41.554	8.171	7.082
	HEM881000244	22.390	13.510	42.662	18.503	18.758	11.192	2.111	13. 188
	HEM881000250	20.878	6.254	20.741	9. 109	1.841	13.561	9.540	2.708
	HEMBB1000258	101.717	75.034	336.781	79. 281	52. 103	67.231	33.313	34. 880
	HEMBB1000264 HEMBB1000266	99.327	57.280 23.082	269.540 23.217	83. 791 14. 456	39.799 28.745	96.654 34.547	62.346 15.022	79. 783 15. 672
45	HEMBB1000272	14.990	14. 502	10.270	6.954	12.730	6.133	4. 205	16.611
45	HEMBB1000274	105. 245	46. 925	190.978	49.759	41.568	43. 127	18.199	25. 826
	HEMBB1000276	6.479	2.218	2.501	4.783	1.754	2.070	2.079	1. 252
	HEMBB1000284	4.790	5.088	7.884	3. 489	2.213	3.213	1.981	3. 304
	HEMB81000307	52.330	30. 191	128.450	28.961	22.039	15.869	9.113	21.677
	HEMBB1000309	86.347	36.463	96.140	43.964	34.442	33.118	18.805	21.507
50	HEM881000312	41.862	30.986	40. 349	24. 933	7.383	79.360	24.114	16.788
	HEMB81000317	49.311	18.053	26. 189	10, 490	10.102	21.107	12.632	13. 384
	HEMB81000318	87. 180	33.847	208. 954	43. 556	23.043	27.764	9, 191	17.641
	HEM881000332	3.892	11.256	14.087	42.331	28.145	14.132	2.408	14.319
	HEMBB1000335	27.939	30.864	21. 167	28.071	12.651	30.027 23.654	12.746	21.753
	HEMB81000336	68.463	26.023 59.290	93. 527	10.608 52.168	22.871 54.197	125.769	23.868 126.562	13. 927 60. 614
55	(HEMOD (00033)	289.853	33. 230	1 33.361	1 32. 100	34, 131	123.103	120.302	1 00.014

Table 41

1	UPSPROTAGAGAA			1.55 1.56				aa 1	00 701
	HEM8B1000338	54.685	45. 765	123.480	44.612	17.722	26.663	17.708	29. 721
	HEMB81000339	144. 258	108, 124	265, 125	105. 421	89.798	89.055	56. 944	50. 241
	HEMB81000341	113.271	46, 622	132, 906	32.751	40.166	37, 986	28.017	30.881
	HEMBB1000343					46.681			
		130.737	71. 935	259.845	80. 183		45. 761	43. 928	46.721
	HEMB81000354	202.146	151. 264_		157.908	153.529	142.579	67. 161	105. 322
	HEMB81000358	92. 244	22.827	29.160	24.670	22.387	48. 989	59. 506	28. 803
	HEMBB1000369	55.720	25. 874	97.758	27.483	21.576	23.750	17.278	16.569
	HEMBB1000373		59.105		61.379	38.792	44. 185	31.504	45. 653
		52. 572		70.779					
	HEMBB1000374	153. 545	115. 183		108. 150	98.073	80.319	58. 214	75. 906
į	HEMBB1000376	95. 394	132.554	369.986	146, 818	60.328	63.876	73.647	43. 202
	HEMBB1000383	37.023	35. 429	24.954	13.017	10.381	22.638	16.842	8.781
						30.891	57.827	37.484	
'	HEMBB1000391	127. 327	30.055	106. 971	24.962				11.921
	HEM881000399	35. 143	10.865	22.406	8.561	4. 100	8. 569	2. 543	8.889
	HEMBB1000402	82.616	20. 485	44.946	25. 430	13.012	19.024	7.725	18.695
	HEMBB1000404	18.903	12.568	10.300	8. 593	9. 455	9. 301	2.672	7.956
	HEMBB1000407	19. 286	8. 572	18. 593	3. 281	2.599	13.454	2.473	3. 407
	HEMBB1000420	95.847	66.573	138.307	54. 950	39. 330	56. 220	37.608	43.081
	(HEMBB1000430	274. 820	161.981	153.601	40.874	406.081	489.107	693.805	115.638
	HEM881000434	350.936	139.4811	599, 497	199. 198	125. 426	113.500	65.776	77.687
	HEMB81000438	67.342	10.187	25.472	7.736	8, 148	27. 875	7.217	6.701
	HEMBB1000441					60. 934			35. 267
		84.086	98. 109	312.643	78. 842		76.141	46. 589	
	HEMBB1000447	76.519	88.156	54.883	26. 628	31. 157	24. 328	25.777	38.008
	HEMBB1000449	22.367	11.282	25. 245	11. 267	1.700	13.053	5. 731	8. 109
	HEMBB1000453	26.781	29.875	49.056	22. 139	35.305	22.456	14.006	15, 902
	HEMBB1000455	37.937	43, 401	129, 423	29, 222	40.584	24. 577	21.227	20.356
	HEMBB1000472	146.390	61, 195	235.753	80. 306.	44. 122	82.882	52.783	87.457
				194, 466			60. 409	34. 897	
	HEMBB1000480	138.135	67.904		46.367	41.944			40.785
	HEMBB1000486	78.511	63.045	211.876	47.786	39.049	36.558	20. 396	17.632
	HEMB81000487	21.510	22.091	29.116	10.718	21.056	15.854	13.086	10.892
	HEMBB1000490	232.419	148, 116	562.064	159.218	134.370	107.861	60. 296	110.306
	HEMBB1000491	149.070	107, 169	349.100	81.342	44, 330	51.147	33.633	59.342
	HEMBB1000492	18. 194	21.930	19.080	9.690	6. 821	10. 632	9. 805	5. 454
	HEMBB1000493	286.390	34. 074	64.876	31.406	23.065	49.816	39. 824	39. 921
	HEMBB1000510	133. 225	95. 239	380.177	165.002	101.728	72. 504	64. 646	83.048
	HEMBB1000516	137.574	35.610	61.963	35. 305	10.932	78.851	39. 905	19. 224
	HEMB81000518	8.388	3.267	26.133	5. 489	1.531	1.500	1.611	1.901
	HEMBB1000523	153.793	88.071	329.880	82.474	43.568	69.756	32. 830	51, 127
	HEMBB1000530	46. 151	13.390	40.950	8.319	32.799	6.126	10.689	8. 426
		62 900							
	HEMBB1000542	57.808	36.831	46. 332	20.306	19,414	5. 489	13. 314	22.747
	HEMBB1000550	39.123	26.036	79.169	22. 945	10.597	23, 147	37. 266	20. 568
	HEMB81000554	192.214	105.635	349.184	148.874	90.632	98.169	55. 377	100.995
	HEMBB1000558	100.759	22.180	68. 289	37.737	35.176	41. 190	47. 163	40.726
	HEMBB1000564	101.412	37, 586	144. 386	37.463	27.344	59.939	31.447	9.452
	HEMBB1000567	361.516	76.515	125.177	66.960	83.698	221.216	145, 840	54. 204
					18, 197		36.942		
	HEMBB1000569	63.847	46.712	54.356		23.752		31. 264	39.479
	HEMBB1000573	99.088	89.487	373.557	76.986	76.236	63.534	39.587	58.648
	HEMB81000575	74.071	67.726	296. 427	63.469	37.530	42. 388	33. 544	46. 151
	HEMB81000579	27.868	12.805	18.934	6.889	3.743	24.452	24. 367	23. 262
	HEMBB1000585	30.826	34, 244	65.882	26.172	19.828	26.184	16.826	33.888
	HEM881000586	85. 397	75.643	187.543	99.762	48, 456	35, 430	28, 693	50.228
	HEMB81000589	135. 404	58.619	243.853	51.181	36. 284	29.883	21.561	27.997
	HEMB81000591	99.680	60. 946	242.306	54. 595	36.589	52.616	32. 332	33.066
	HEMBB1000592	30.320	18.740	34. 338	11.753	8.732	28.305	13.707	12.164
	HEMB81000593	148.639	68.816	255.892	61.084	46.829	61.565	49. 545	66.588
	HEM881000595	27.140	21.001	29.869	21.272	9, 199	21.841	15.487	29, 680
	HEMBB1000598	39.074	31.891	85.011	22.815	13.772	21.958	13, 576	26.747
	HEMB81000611	14. 828	6.552	11.601	7, 498	7.461	15. 614	9. 246	9. 161
	HEMBB1000617	193.986	137.945	458.678		87.855	84. 583	46. 273	77.986
	HEMB81000623	65. 566	26.480	50.777		18.923	40. 974	28. 571	23. 219
	HEMBB1000630	62.606	23.074	40.815	18.796	14.186	31.973	21.492	13.779
	HEMBB1000631	61.311	41.283	27.586	23, 498	24.433	35.043	48, 566	22.826
	HEMB81000632	58.747	55. 433	156.750	30, 460	29.661	33.497	21, 899	21.857
	HEMB81000636	127. 885	47.562	59.456	48. 965	33.643	65. 366	42.360	37. 349
	HEMBB1000637	817.391	628.017	1645.738	524.605	482. 307	443.855	191.753	265. 704

Table 42

	HEM8B1000638	55.058	47.453	95.751	42. 262	25.684	15.056	22. 121	28.829
	HEMBB1000642	179. 188	88.317	251.754	80.865	42.468	81.296	37.696	52.009
_	HEMBB1000643	43, 411	25.689	113.037	18.985	11.038	14. 245	8. 276	18.743
5									
	HEMBB1000649	27.852	45. 202	137.371	34.816	24. 496	9. 967	11.881	22.322
	HEMBB1000652	84.942	61.856	126.562	78. 131	42.090	36. 343	22.852	31.597
	HEMBB1000655	418.308	73. 377	56.858	57. 166	32.733	57.424	38.897	44. 477
	HEMBB1000665	16.253	13.954	10.766	20.817	6.796	13.110	7. 987	4. 458
	HEMBB1000668	28. 587	13. 435	14.606	13.788	25. 844	15.049	12. 549	11. 202
	HEMBB1000671	239.020	122. 952	561.221	119.970	96. 244	75.058	66.812	88.267
10							4.811		
	HEMBB1000673	11.633	5. 779	14.629	14. 904	5. 916		2. 141	12.812
	HEMB81000679	16.899	7.357	23.438	7.697	1.049	30. 246	7.774	7.063
				430.254		66.411	89.722	49, 173	67. 832
	HEMB81000684	188.240	157.754		128. 150				
	HEMBB1000692	4.978	9. 265	11.569	5.085	1, 158	3. 240	3. 421	1.785
				59, 522	22. 326	25. 408	13.898	31.488	20.706
	HEMBB1000693	63.119	40. 561						
45	HEMBB1000705	15. 560	31,798	122.757	36. 451	19.928	11.568	2.839	10.179
15	HEMB81000706	22.553	13.626	23.777	8.621	11.683	41.509	10.019	7.584
	HEMB81000709	74.737	77.864	245.726	50.833	51.093	50. 427	37. 955	51.357
	HEMB81000714	23.726	10.733	6.625	12.298	6.349	9.891	2. 142	14. 350
	HEMB81000725	24.239	9. 575,	11.437	13.761	12.596	17. 372	8. 105	16. 144
	HEM881000726	86.971	84. 395	208.396	65. 157	43, 881	37. 441	22.020	39.067
20	HEMBB1000729	51.556	25. 288	140.931	23.005	27.775	18.629	12.838	14. 902
20	HEM8B1000738	39.002	38. 955	166.616	42, 588	21.380	43, 330	7.181	21. 192
	HEMBB1000749	115.917	94. 942	454.741	136.454	54. 340	39. 253	32. 933	49. 141
	HEM881000763	47.835	25, 201	36.488	16. 952	21.036	31.919	14.990	12.111
	HEMBB1000770	30. 598	45. 410	167.003	32.786	26.482	25. 698	18. 186	24. 127
		30. 538							
	HEMBB1000774	27.168	21.690	33.470	20.937	12.916	22. 598	8. 092	17.606
	HEM881000777	246.286	57. 131	58,743	31.851	40.345	119.113	81.364	53.990
25					<u> </u>				
20	HEM881000781	41.945	36.620	34.149	24. 543	23.561	16. 383	14. 371	20.775
	HEMB81000788	10.756	10.608	5.481	5.429	2.950	5. 995	4. 522	4. 589
							16. 477	7.916	7.672
	HEM881000789	28. 490	9. 620	26. 151	16.088	11.640			
	HEMBB1000790	74.318	56. 925	185. 959	63.749	33.523	24. 232	24. 414	28. 423
	HEMB81000794	18.080	17.254	38.876	24. 305	7.427	10.338	5. 445	9. 305
	HEMBB 1000807	50,070	31.869	22.751	19.865	20.934	27. 002	18. 350	27. 280
30	HEMBB1000809	334.541	42.976	42.300	26.454	9.545	31. 526	31.677	44. 152
	HEMB81000810	189.365	50, 676	163.325	33.349	38.994	74. 400	45. 398	19. 262
	HEMB81000821	40.710	9.304	21.006	6.841	5. 422	15. 981	10.835	5.685
	HEMBB1000822	8.726	3. 570	3.541	1.411	7. 255	5.519	1. 285	1.525
	HEMBB1000826	68. 485	40. 348	201.149	68.467	43. 204	31.769	32.812	55. 367
	HEM8B1000827	50.671	34, 326	108.391	32.945	15.076	25. 813	18.713	25. 457
35	HEMBB1000831	38.060	20.466	29.131	12.368	19.990	20. 562	25. 373	6.415
	HEMB81000835	59. 181	56.345	127.358	58, 150	44.350	35.831	25. 687	35. 108
	HEMB81000840	117.639	63. 375	340.802	61.186	48. 924	38. 995	20.712	30. 526
	HEM881000848	98, 938	53.024	210.423	42.569	28.984	47.603	29.642	29. 431
				0.621		1.621	1.272	1.364	1.086
	HEMBB1000852	1.827	2.160		2.559				
	HEMBB1000857	16.897	16.768	19.951	14.921	12.912	17. 270	10.179	14.915
40	HEMBB1000858	25.634	16. 531	8. 162	8.209	14, 482	12,749	92. 823	10. 102
40									
	HEMBB1000867	106.946	56. 331	264.748	50.278	36.949	41. 202	26.795	29.760
	HEM881000870	68.550	62. 423	192.351	52.406	39.303	55. 641	23.738	27.427
						7.689	10.690	11.143	26.241
	HEMB81000876	21.813	12.044	24.968	11.314				
	HEMB81000881	30.089	16.478	28. 345	14. 926	18.419	17. 763	18. 901	20.494
	HEMBB1000883	11.669	10. 263	26. 185	6.975	2.780	8. 223	2.906	3.540
								44. 444	2. 340
45	HEMBB1000887	42.638	32. 274	66.780	22.979	31.512	42.842	20. 622	22. 566
	HEMBB1000888	20.318	8. 193	11.483	5.178	4.073	8.708	6.801	4. 342
	HEMBB 1000890	40.795	42. 287	112.076	25.031	11. 171	23.116	15. 491	16.447
	HEMBB1000893	38. 227	10,603	88. 306	24. 535	14.440	12.863	9.734	17.727
	HEMBB1000900	23.814	8.709	17.013	9. 267	10.928	12.199	14. 105	11.108
	HEMB81000905	63, 589	43, 501	37, 125	41.367	26, 379	29.649	38.699	31.891
						28.838	28.194	15.897	26.230
50	HEMB81000908	42.944	54.674	120.821	34. 982				
	HEMB81000910	72.960	51.795	161.850	41.050	36.594	37.378	13.612	23.263
						12.371	26.067	14.715	19. 258
	HEMB81000913	33.820	35. 219	96. 448					
	HEMB81000915	1910.513	222. 511	693. 345	124.825	532.993	1548. 228	1159.943	223. 176
	HEMBB1000917	99.638	64. 212	310.142	53.316	39.091	34. 989	22. 324	40.667
	HEMBB1000927	80.569	11.252			21.944	24. 546	17.769	17. 391
	ILICHOO LAAAAAA	1 44 130	33. 556	95.029	29.041	17.945	21.758	22.973	31.034
55	ILE DE LUMBAY	1 33 1/5							
55	HEMB81000932	33.128	1 33. 330	1 30.023	1 60.041				

Table 43

	(UEVED LAGRAGA	000 000	300 03r I	COC 050	000 F41	111 660	530 431 I	202 100	291.706
	HEMBB1000933	883.639	393.035		289. 543	312.660	538, 431	353. 155	
	HEMBB1000936	23.212	17.243	46.380	14. 205	25. 527	13.908	8. 530	11.716
5	HEMBB1000939	105.016	36.905	52.525	19. 304	30. 111	35. 223	41.856	37. 146
	HEMBB1000941	6. 540	27. 555	15. 872	4.660	6.130	17.648	83. 246	9. 541
	HEMB81000947	36. 384	18.020	47.143	21. 361	9. 565	34. 299	13. 482	13. 269
	HEMBB1000954	16.970	17.878	19.810	11.407	6.851	17. 302	10.023	8. 877
	HEMB81000959	22. 223	21. 226	78.296	22. 443	5. 599	10.268	10.673	12. 183
	HEMBB1000973	11.584	10.364	21.189	8. 579	7. 102	23.845	5. 510	9, 891
10	HEMB81000975	99.598	37.022	69.027	23.084	27.137	40. 162	56. 997	30.316
10	HEM881000981	10.199	12.524	23.602	20. 141	5.813	6. 152	13.771	4. 102
	HEM881000985	13.065	8.026	7.574	4.776	6.642	2. 985	6.049	3.612
	HEMBB1000991	67. 124	17.092	28.053	8.864	8. 560	28. 394	25.072	10. 907
			127.636	352.650	90.350	64. 926	71.240	60.014	102. 622
	HEMBB1000996	170. 256			15. 226	10.611	14. 541	7. 598	9. 642
	HEMBB1001000	48. 257	19.380	16.573		0.000		0.000	
15	HEMBB1001004	0.797	1.839	0.439	0.000		0.318		0.000
	HEMB81001008	17. 533	13.975	16.434	11. 194	6. 400	12. 238	6.478	9. 235
	HEMB81001011	39.743	19.337	28.396	15.752	15. 302	17.720	15. 586	16.702
	HEMBB1001014	121.726	46.352	244.715	50.619	33.004	55.708	30.100	34. 469
	HEMB81001020	86.065	68.022	243.352	67.763	53. 522	50. 406	30. 247	49. 844
	HEMB81001024	66.546	59.010	205. 347	41. 480	31.865	35. 052	21.045	39. 489
20	HEMB81001026	36.265	27.027	76.443	19.990	25. 484	27.657	12.014	23. 129
	HEMB81001037	64.392	37.810	120.090	20, 652	22. 459	27. 294	18.918	28. 917
	HEMBB1001042	58. 936	20.428	42.468	17. 255	15.600	32.463	20. 274	18. 506
	HEMBB1001046	76.790	22.021	40.791	13.932	17.825	47.853	26.672	30.056
	HEMB81001047	76.665	39. 237	208.757	53.469	44. 539	37.624	16.049	20. 262
	HEMBB1001048	133.028	58. 176	140.515	48.390	34.614	42. 111	29. 526	34. 858
25	HEMB81001051	22.699	8.465	13.142	9.942	10.065	9. 946	5. 881	8. 790
	HEMB81001056	40.040	16.494	45,000	22.674	18.685	21.131	18. 431	12.498
	HEMBB1001058	88.873	59.116	223.822	45. 122	34. 696	29. 783	21.562	25. 222
	HEMBB1001060	35. 486	18.631	33.852	60.851	26.807	13. 499	12.993	19. 391
	HEMBB1001063	53. 418	36.359	125.166	33. 156	24. 220	19. 182	16. 188	14. 597
	HEMB81001068	79.181	46.879	78.756	35.034	26.835	79.006	63. 198	43. 296
30	HEMBB1001082	66. 296	58. 491	173.393	49.675	25. 253	33.015	14. 189	22.904
30 .	HEMB81001095	64. 435	31.409	20.825	17.116	14.939	41. 581	21.497	13.792
	HEMB81001096	43. 372	28.562	94. 366	32.120	13.089	21.236	15.814	22.034
	HEMB81001101	79. 652	21.131	40.775	18.757	35.350	46.263	18.855	13.874
	HEMBB1001102	51.740	27. 585	86.794	21.160	12.958	16.450	7. 235	8. 605
	HEMB81001104	61.846	33. 489	28.997	14. 789	10.623	20.859	15. 993	10.658
0.5	HEMBB1001105	69.199	32.868	132.855	27. 292	32.605	49.984	20.779	23.751
35	HEMBB1001112	161.356	78.361	73.588	64.617	86.150	93.363	87.696	95. 854
	HEMB81001113	114.744	130.208	298.139	107.218	73.757	61.718	32.824	66. 952
	HEMB81001114	105. 358	95.960	365.719	66.457	62.314	35. 251	34. 480	51.970
	HEM881001115	67.274	16.815	13.190	26.838	17.638	29. 948	23.803	34. 239
	HEM8B1001117	2. 434	10.619	14. 951	4. 152	4. 937	2. 694	2.729	18. 952
	HEMB81001119	18.198	17.501	58.077	15.560	5. 202	13. 437	5. 261	9.614
40	HEMB81001126	306.301	111.345	266.365	81.302	76, 905	130.782	58. 863	61.487
	HEMB81001133	39, 673	35.703	178. 312	45. 328	36.363	38.712	14. 400	26.997
	HEMBB1001137	53. 424	19.209	46.849	14. 453	13.705	30.395	18.865	15.761
	HEMBB1001142	105. 888	131.411	405.403	98.008	104.700	62.754	32.598	75. 485
	HEMB81001145	114. 864	106.329	348.161	78.364	57. 587	54. 983	24.738	51.568
	HEM881001151	149.618	23.632	66.607	14.582	34. 238	68.060	46.084	19.806
45	HEMBB1001153	92. 263	53.444	153.351	44. 131	37. 191	34.991	21.708	32. 599
	HEMBB1001158	64. 416	30.844	50.578	22.880	32. 523	47.046	24. 553	39.658
	HEMB81001169	95. 424	70.158	253.814	76.490	44.058	37.113	24. 102	38.757
	HEMBB1001170	34. 989	7.730	32.617	5, 324	4.217	11.418	7.623	5. 208
	HEMBB1001175	46. 512	27.401	45. 252	21.001	15.416	20.636	17. 361	36.021
	HEMB81001177	126. 389	86.212	396.633	84. 357	48.470	40.910	34. 438	42.680
50	HEM881001182	70.825	30.508	45.077	19.262	28. 316	32.507	25.771	26. 488
	HEM8B1001192	30.059	21.703	61.610	20.151	5.688	22.456	24. 299	31.214
	HEMBB1001199	1.469	0.000	0.000	4.430	0.797	2.148	1.260	1. 223
	HEMB81001200	2. 266	1. 426	2.071	5.734	0.000	2.413	1.567	2.969
	HEMBB1001208	111.969	37.738	122.154	28. 426	28.653	55. 253	32.443	21.624
	HEMBB1001209	103.602	77.445	233.649	60.849	25.456	40.993	26.273	33.636
55	HEMBB1001210	14. 499	40.527	32.902	6.231	10.125	16.413	17. 251	28.930

Table 44

	HEMBB1001215	219, 922	83.033	126. 326	63.007	71,733	115. 441	61.961	72, 230
	HEMBB1001217	63, 633	22.116	41.047	17.479	20, 160	53.164	31.645	18.739
	HEMBB1001218				53. 412	29. 467	23.819	20. 495	24.079
5		98. 226	47.137	142. 266					
	HEMBB1001221	0. 524	1.310	12. 795	0.988	0.992	0.867	0.000	1.767
	HEMBB1001224	52. 109	37. 281	86.318	28. 364	24. 177	19.072	16. 478	20. 321
	HEMBB1001230	38. 785	17.158	30.714	15. 256	12.698	31.469	27. 596	17. 436
	HEMB81001234	335. 966	64.817	131.669	43.601	69. 385	167. 134	101, 415	57. 258
	HEMBB1001235	152.870	67.952	84. 726	40. 262	26.665	52.686	38. 623	49.693
10	HEM8B1001237	16, 971	23.623	33.663	30.744	21.161	18.495	18. 264	25.643
10	HEMBB1001242	26.787	15,776	22.922	4. 200	5. 187	11.277	10.621	7. 589
	HEMB81001244	280. 439	9. 589	9.743	8. 128	2.116	4. 366	2.735	2. 871
	HEMB81001249	51, 892	27.766	106.010	25. 983	19.890	21.254	16, 839	21.542
	HEMBB1001253	50. 869	33.773	58. 857	31.656	8. 253	38. 144	20.639	25. 942
						6. 376	18.461	22. 558	
	HEMBB1001254	28. 109	8.716	61.080	12.779				8. 559
15	HEMBB1001266	2.010	9.088	3. 704	1.682	16.420	18.653	1.717	1.611
	HEMBB1001267	131. 334	93.697	391.730	88.886	45.610	62.418	33. 457	63.350
	HEMB81001271	31, 480	28.408	63.773	19.821	15. 244	12. 530	8, 683	10.739
	HEMB81001282	41.156	11, 440	25. 546	10.847	7. 531	21.762	15. 737	10. 592
	HEMB81001287	195. 274	200.678	131.870	63.454	15. 491	70.758	43.360	52.931
	HEMB81001288	40. 232	10.227	25. 481	9.789	5. 520	21.519	16.538	9.861
20	HEMBB1001289	84. 233	74.730	246.417	61.615	31.689	36.447	24. 521	38.077
20	HEMBB1001290	57.742	13.181	11.174	33.921	23. 320	24.860	82.615	15. 369
	HEMB81001294	80.761	23.745	72.937	16.689	20.147	45. 268	37.686	22.951
	HEMB81001299	58.616	17.094	44. 424	13.532	14.650	31.325	32, 822	12.329
	HEMB81001302	87, 107	24.979	56. 357	23.389	20.784	37. 921	28. 849	21.981
	HEMBB1001304	12, 134	0.119	5. 246	19.403	1.810	3. 978	2.153	1.580
	HEMBB1001314	6, 410	5. 111	25. 042	5. 961	3. 244	7.037	2. 954	2. 258
25	HEMB81001315	3, 706	8.398	10.733	3.067	1.405	3.652	1.659	1.943
	HEMBB1001317	39, 137	34.918	87.084	32. 290	25. 473	21. 551	14.009	18.118
	HEMBB1001326	13. 902	5.726	7.704	2.886	2.324	1.546	2.008	5.612
	HEMBB1001331	34. 871	17.866	37. 859	11.626	6. 188	23. 138	24, 975	17. 786
						15.964			
	HEMBB1001335	22. 550	20.911	19.341 187.675	12.458	52.185	18. 477	15. 941 25. 750	5. 614 29. 233
30	HEMBB 1001337	61. 645	43.894		45. 250				
00	HEMBB1001339	20.634	25.030	21.230	11.541	12.874	18. 490	12.601	13.466
	HEMBB1001344	31. 209	8. 322	15.710	5.412	6.749	16.517	16. 482	9. 869
	HEMBB1001346	44. 149	21.512	38. 191	15, 415	9. 432	26.936	17.706	15, 965
	HEMBB1001348	66. 624	40.319	173. 356	39.887	26.835	31.783	20.641	26.670
	HEMBB1001350	103.603	17.400	35, 832	13.555	13.837	54. 503	34, 694	19. 925
	HEMB81001356	12.440	11.385	25. 095	8. 592	6.787	7.806	8.759	8. 923
35	HEMB81001364	28. 525	14. 483	31. 452	11.829	13.494	12.620	13, 025	10.117
	HEMBB1001366	57. 883	53.690	210. 263	52. 112	27. 208	41.191	29. 156	32.064
	HEMBB1001367	140.660	59.744	283. 101	54. 260	46.338	67.368	43. 944	48. 485
	HEMBB1001369	17. 341	20.708	71.044	14.855	7.629	12.537	7. 158	14. 407
	HEMBB1001380	50. 204	67.647	124. 463	41.290	43.730	41.591	29.026	63.358
	HEMBB1001381	19.588	19.545	34. 218	14.113	18.710	9. 428	10. 202	13.801
40	HEMBB1001384	17,779	11.154	26. 926	11.606	19.030	10.038	7. 367	14, 535
	HEMBB1001387	20.705	16.837	19.148	9. 955	8.901	15. 994	7. 831	13. 345
	HEMB81001394	21.419	19.091	32.720	17.551	19.172	11.590	12.282	11.322
	HEMBB1001407	39, 158	17.718	75.721	24.299	17.481	17.410	20. 342	15. 525
	HEMBB1001410	18, 880	3.346	6.042	2.907	2.655	0.000	2.839	2.094
	HEMBB1001413	32. 291	25.769	80.279	17.033	21.102	11.132	12.610	24, 207
45	HEMBB1001419	36, 323	42.415	185. 239	24.790	21.849	17.972	13.895	31.342
10	HEMB81001421	29. 464	57.495	109. 370	12.065	15.685	64. 181	165.647	23. 322
	HEMBB1001424	9.663	7.148	10. 294	6.073	6.773	7.183	5. 215	8. 524
	HEMB81001426	36. 471	25.897	86.872	20. 138	17.823	19.534	15. 347	23.782
	HEMB81001429	60. 351	47.669	39. 928	29.802	21.695	39.456	39.474	41.210
	HEMBB1001436	168. 445	86.814	350. 902	88.825	54. 545	86.724	48. 813	58. 527
50	HEMBB1001443	20.733	11.137	12. 445	8.769	16.707	14. 531	9. 581	12.477
	HEM8B1001449	70. 239	34.064	146.511	28.311	23.391	19.979	16.080	22.377
	HEM8B1001454	60.851	40.766	133.878	33.168	28.709	36.541	29.720	26.623
	HEMB81001458	77. 938	28.808	33. 472	15.970	29. 260	40. 965	25. 268	28.079
	HEMB81001461	44. 192	44. 580	179. 531	65.974	16.217	45. 935	14.669	27.974
	HEMBB1001463	57. 949	102.937	230.980		41.957	48. 857	25. 233	38.517
55	HEMBB1001464	18.058	9, 999	14. 908	10.039	7. 528	8. 580	2.638	2.964

Table 45

	HEMBB1001466	31.340	22.324	20.480	15.496	3.611	15. 533	10.020	13.761
	HEMBB1001482	12,741	4,057	9. 987	4. 175	4, 887	24. 039	4.114	4. 470
			21.417	65, 107	17.492	9, 196	12. 958	8.167	14.603
5	HEMBB1001500	26.823							
	HEMB81001505	116.783	105. 297	302.199	104. 682	36.419	54. 346	38. 027	46. 591
	HEM881001521	55. 379	38.602	133, 188	25.792	20, 204	23. 504	18. 528	22.786
	HEMBB1001527	331.186	160.160	252. 225	131.308	116.694	179. 333	72.732	79.869
	HEMBB1001530	24.722	25.693	57.090	19.457	7.662	20. 875	31.031	23.503
	HEMBB1001531	43.913	51.679	130. 225	34.674	21,061	27.704	18.966	32.578
	HEMBB1001532	6. 957	3.901	34. 322	7.593	1.875	8. 172	300.808	7.501
10									
	HEMBB1001535	71.654	59. 202	131.794	46.369	28. 936	34.644	21.690	23.017
	HEM881001536	73. 109	48. 204	106.813	35. 175	16.411	22.356	19. 126	20. 785
	HEM8B1001537	40, 809	54.756	140.043	43, 830	21.583	31. 273	8.692	29. 500
	HEMBB1001542	79.436	33.152	94. 294	34. 360	26, 100	44. 300	19.679	22.657
	HEMBB1001543	55.819	14. 588	8. 417	4. 239	7. 702	20.740	11.834	18.032
15	HEMBB1001547	10.746	8. 433	12.415	9, 202	10, 101	15.047	10.631	8.198
10	HEMBB1001548	163.125	42.223	39.134	33.781	26.421	115.789	76, 174	67, 211
	HEMB81001551	32.248	10.176	8. 937	9.728	20.037	69. 247	7078.074	11. 439
	(HEMBB1001555	62.998	58.959	166.842	57.865	40.731	30. 981	17. 189	40.721
	HEMBB1001562	67.088	35. 544:	83. 929	24. 475	18. 852	28. 472	27.682	23. 295
	HEMBB1001564		320. 422	580, 390	304.052	124.857	300.720	202.502	439. 361
		139.467							
20	HEMBB1001565	56.749	43. 545	123.727	39.891	29. 530	30, 029	17.527	28. 501
	HEMB81001569	34, 482	26.904	100.487	28.883	16. 462	19.020	8.403	16.605
	HEMBB1001573	48, 940	40.308	65. 598	41.979	32. 247	35. 238	25. 583	36.979
	HEM8B1001585	153. 364	57.831	211.685	61.076	40.832	38, 446	18.915	42.636
	HEMBB1001586	44.946	40.343	113. 224	34. 426	18. 386	24.673	16.535	26.124
	HEMB81001588	157.947	130, 811	402,650	111.293	69, 831	80.240	46.050	75. 499
	HEMBB1001595	12.602	11.160	44. 464	13.949	6.811	11.538	4. 359	11.569
25									
	HEMBB1001596	53.986	20.798	39.629	25. 473	20.578	32.621	23. 309	36.564
	HEMBB1001599	29. 275	7.352	13.267	11.568	5. 279	15.756	10.260	5. 135
	HEMBB1001603	3.581	2.642	7.782	4, 279	3.051	0.341	1.424	3, 160
	HEMBB1001606			7. 226	7.657	3. 104	5. 383		4. 364
		6.897	7. 220					5. 658	
	HEMBB1001612	101.576	58.128	240.469	58.770	36. 287	42.917	27.221	40.063
	HEMBB1001618	52.604	38.648	141.745	37.723	24. 274	24. 922	17.197	24. 223
30	HEMBB1001619	59. 431	78.268	138, 545	63. 285	52. 275	37.035	22. 185	38.081
	HEMBB1001623	33.128	8.489	11.122	6.318	8. 326	16.007	3. 331	7. 918
	HEMB81001625	10.068	16.076	8.496	7.577	2. 293	8. 389	1.716	4. 647
	HEMB81001630	7, 144	5. 464	31, 186	8.383	3. 256	11, 196	3.053	5.942
	HEM881001635	18.151	8. 186	33.138	13.501	9, 143	9.688	44.037	8.859
								20.792	
35	HEMBB1001637	40.224	35. 174	58.964	24.082	26.640	26.340		26.243
35	HEMBB1001641	21.655	10.768	33.553	9. 122	5.845	7.210	5.796	8.300
	HEM881001653	76.458	45. 984	138. 114	33.606	30.023	33, 136	16.720	25.949
	HEMBB1001665	3.000	0.352	5.654	0.275	0.718	0.106	0.899	0.407
	HEMB81001666				22.201	9.196	20. 512	10.659	15.687
		48.027	23. 276	59.669					
	HEMB81001667	2.570	7. 909	3. 107	5.847	8.690	2.748	1.999	8.738
	HEMB81001668	2.545	8.885	13.392	8.498	18.131	3. 355	1,531	3.932
40	HEMBB1001669	5.751	5. 364	10.395	3.219	4. 970	5, 110	4.341	2.139
								12.909	
	HEMB81001670	17.795	10. 903	34.891	20.715	11.725	22.401		20.514
	HEMB81001673	69.924	44. 194	58.806	53.036	21.640	40. 433	25.038	49. 339
	HEM881001675	58, 961	13,650	21.648	10.914	9.356	22, 270	15.894	11.977
	HEMB81001679	51, 245	9. 166	29.461	6.718	11, 101	24.642	13.266	4. 383
	77.4				1	 	2.4		12 22 1
	HEMBB1001684	27.854	11.218	30, 139	14.655	11.546	25.422	15.072	13.683
45	HEMBB1001685	9.626	8.721	34.446	7.134	4.659	1.316	3. 180	6.172
	HEMBB1001695	2.706	4.723	4,741	1.162	8.059	1.109	1.036	1.119
	HEMBB1001703	116.774	37.756	115.693	36.901	34. 790	69. 383	44. 901	43.576
	HEM881001704	67. 385	52.606] 211. 228	52. 452	40. 405	43.432	33.952	54.662
	HEMBB1001706	122. 282	70.475	227.746	77.627	63.608	53.010	38.740	56.789
	HEMBB1001707	111.416	69.815	154. 286	51.656	60.773	50. 260	33.306	43.746
50	HEMB81001717	14.112	16. 260	60.454	10.609	5. 688	9, 921	4.816	8.073
	HEMBB1001731	29.550	36. 222	21.992	33.872	22. 551	35.654	37.976	32.089
	HEMBB1001734	75.818	39.477	107.419	26.507	15.856	20.715	17.010	17.320
	HEMBB1001735	63. 245	22.136	169.823	34. 289	26.478	18.371	17.292	27. 924
	HEMB81001736	20.722	18.061	27.944	17.598	12.534	9. 551	10.504	13.178
	HEM881001747	21. 158	15. 281	18.501	9.967	9.806	11.088	17.268	12.572
55	HEMBB1001749	89. 421	90. 342	429. 206	126. 585	53.728	56.733	28.560	64.467
JJ	(IICM001001143	1 03.421	30.342	743.200	1120. 383	1 33.120	1 30.133	1 20.300	07.401

Table 46

	HEMBB1001753	85. 135	63.020	101.881	44.766	60.100	46.138	48. 988	44. 990
	HEMBB1001756	86. 556	37.048	83. 531	33. 276	42.763		32.005	30. 821
							54. 273		
5	HEMBB1001757	1. 981	3. 522	5. 232	3. 590	1.394	7.486	3. 256	3.014
•	HEMB81001760	13. 573	14.554	27.053	7. 204	5. 280	8.129	5. 242	4. 088
	HEMB81001762	26. 210	15.945	24.826	8. 467	6.461	26. 934	6. 893	9.656
	HEMBB1001780	18.738	33, 363	27.562	17. 311	13.893	4, 277	14. 584	19. 429
	HEMB81001785	3.266	2.954	7.974	3. 522	3.900	7,429	3.964	4.008
	HEMB81001788	77.710	51.716	232.298	72.096	40.555	41.418	29. 586	33. 423
	HEMB81001793	221. 348	29.215	45. 528	20. 500	22. 918	33. 927	36.095	
10									25. 245
	HEMB81001797	4.049	9.015	10.442	4.015	2.532	8.773	2. 904	6. 333
	HEMB81001802	430. 563	24. 213	34.832	14. 183	17. 392	26.448	23. 001	29.744
	HEMBB1001812	91.804	71.389	218.174	56.457	56.645	54.459	15.772	55. 255
	HEMB81001815	506.853	426.652	275. 995	120.005	129. 468	289.852	148.011	122. 368
	HEMBB1001816	90. 696	55. 478	178.334	52.637	25. 170	45, 331	35. 194	47. 899
	HEMB81001831	22.874	14. 551	46.474	16. 825	9.329	19. 975	9.745	18.634
15	HEMBB1001834	456.615	299.793	406. 927	241. 148	284. 283	499. 103	267. 485	306.611
	HEMB81001836	138. 292	91.469	348. 309	101. 544	73.058	67.103	40. 539	76. 261
	HEMB81001839	9. 720	6.600	7.318	0.000	2.606	4. 296	2. 217	2.738
	HEMBB1001841	345. 524	134.230:	67.049	25. 938	60.560	21.530	21. 177	18. 486
	HEMBB1001844	61.041	25. 820	34.819	14. 237	14.648	34. 333	20.655	31. 102
	HEMBB1001847	126.241	111.341	239.722	147.873	65.849	86.164	47. 980	108. 378
20	HEMB81001848	40. 802	39.856	24.837	12.646	9.727	18.893	18.093	17. 754
	HEMB81001850	171, 151	101, 141	118.680	33. 622	64.050	118.364	50. 599	75. 857
	HEMBB1001859	133.676	77.853	231.163	65.024	41.660	123, 173	103. 961	48. 695
	HEMBB1001863	115.353	92.421	255. 141	83. 601	85.833	53.693		
								30. 832	49. 888
	HEMBB1001867	15. 427	15.822	8.336	10.061	4.673	8, 415	6. 299	9. 816
0.5	HEMBB1001868	24.470	17.457	24. 238	7.996	8.810	8.133	10. 520	11. 923
25	HEMBB1001869	82.894	76.711	234. 322	61.007	44. 801	45. 547	29.853	39.008
	HEMBB1001872	15. 921	7. 288	5. 998	10. 151	2. 561	5.674	9. 542	5. 964
	HEMB81001874	36.336	11.065	22.113	15. 221	9.515	14.138	6.058	5. 891
	HEMBB1001875	7.615	19.234	13.755	26. 314	11.646	3.662	5, 863	7. 228
	HEMBB1001880	107.638	82.806	115.014	59. 163	39.712	47.440	27. 454	37. 214
	HEMB81001899	15.785	11.630	15.181	7, 571	2. 259	12.203	4. 190	3. 366
30	HEMB81001903	59.215	24.149	27. 564	15. 205	8.601	28.805	15. 592	15. 765
••	HEMB81001905				15. 117	8. 559		12.021	
		29. 932	24.402	20. 256			17.138		12.009
	HEMB81001906	15. 456	13.077	51.260	10. 147	16. 547	10.906	7. 943	9. 129
	HEMBB1001908	35.095	32.316	100.465	26. 514	24.742	20.649	8. 759	14. 223
	HEM881001910	67.419	35. 922	139.126	58. 266	43, 100	26.178	19. 330	29.710
	HEMB81001911	50. 456	46.682	196.311	58. 337	31.782	35. 278	19. 934	32.009
35	HEMB81001915	40.796	27.017	19.351	20. 885	15.345	12.662	9. 798	36. 052
	HEMB81001921	95. 398	115.190	314, 157	85.049	59. 940	59.397	36.034	60. 585
	HEMB81001922	54. 587	37.299	107.814	29.796	15,712	23.741	15.662	16.568
	HEMBB1001925	35. 478	39.156	106.631	23. 241	15.055	16.405	13. 936	15. 471
	HEMB81001930	9. 272	7.467	11.545	7.045	3. 402	5.636	2. 969	5. 808
	HEMB81001944	122. 259	83. 163	268. 572	86. 582		51.236	27. 262	45. 542
						66.995			
40	HEMB81001945	55.555	20.668	28. 702	7.169	21.076	24. 208	18.042	10. 472
	HEMB81001947	47.254	12. 987	21.887	16.223	6. 133	25.673	16.697	13. 440
	HEMB81001950	99.345	31,711	42. 202	32.724	17. 168	68.211	28.763	30. 429
	HEM881001952	67.117	40.169	164.691	39. 168	16. 287	31.103	11.276	24. 511
	HEMB81001953	56.049	47. 572	147.635	34.659	22.662	21.660	13.445	22. 280
	HEMBB1001957	43.669	20.350	106. 261	26.369	16.837	16.589	5. 199	12.837
AE.	HEMBB1001959	26.731	45.573	72.402	48.003	21.477	24. 554	17.194	36. 361
45	HEMB81001962	59.585	38.413	125.747	48.471	52.786	46.598	20.834	29. 320
	HEM8B1001967	156. 252	96.306	460, 639	121.361	89.090	70.066	46.606	68. 839
	HEMB81001973	62.418	55.111	203. 353	61.777	40, 564	39. 531	24. 193	43. 482
	HEMB81001978	205.611	67.998	184.804	55.506	42. 195	56.711	62.043	55. 171
	HEMB81001983	115. 219	97.908	189.950	79.417	69.496	62.957	41.995	65. 291
50	HEMB81001987	23.094	30.009	63.743	16.838	10.970	10.414	5. 543	10.645
	HEMB81001988	26.549	17.876	71.399	12.651	11.631	11.873	6.563	10, 248
	HEMB81001990	61.049	28.808	125.791	31.477	30.752	26.525	9.894	24. 366
	HEMB81001996	40.435	12.303	17.096	14. 159	3.837	18.573	11.696	13. 433
	HEMB81001997	91.453	62.313	247.838	64.724	40. 131	29. 522	27.492	42. 942
	HEMB81001999	28. 583	9.839	33.748	34. 520	11, 455	23.048	14.798	
									25. 158
55	HEMBB1002002	19. 354	10.115	14. 415	9. 527	16. 781	12.044	7. 088	14. 724

Table 47

	TUCTION LONGOC	107 100							
	HEMBB1002005	127. 202	87.407	314.165	82.406	66.505	55. 577	40.792	64. 185
	HEMB81002009	0.000	1.364	22.770	0.807	4. 369	1.295	0.000	0.000
	HEMBB1002013	28.258	13.676	16.813	10.399	10.765			
5							17.046	7.782	9. 691
	HEMBB1002015	105. 576	48. 524	66.937	36.377	38. 220	74.637	28. 221	34.621
	HEMBB1002024	216.724	27. 841	16.159	12.961	10.268	16, 725	13.378	30. 580
	HEMBB1002035	46.139	20, 267	93.090	25.830	19.155	14. 290	9.089	
	HEMB81002039								10.861
		56.819	33.510	91.779	23.686	12.816	13. 451	13.710	16.666
	HEMBB1002041	64.639	34. 426	51.061	22.611	27. 241	31.364	25. 209	28. 240
	HEMB81002042	108.989	70. 262	244.087	61.596	54.097	58. 195	45. 407	53, 478
10	HEMBB1002043	45.022	36.752	179.777	48. 242				
						21.779	25.603	30.919	28. 446
	HEMBB1002044	13. 181	2.012	5. 797	1.053	1. 982	1.313	3.432	2.045
	HEMB81002045	289.530	197.322	441.790	143. 182	150.349	206.083	108.290	118.515
	HEMBB1002049	35.193	24. 481	83.015	26.999	19.710	27. 535	16.278	24. 921
	HEMBB1002050	37.095							
			16. 954	49.110	12.868	13, 580	16.690	9. 422	14. 540
15	HEMB81002051	36.389	19.655	68. 218	18.665	8.800	22. 352	16.403	17.616
75	HEMBB1002068	75. 935	30. 174	53. 312	27.588	23.758	28. 553	40. 522	36.664
	HEMBB1002069	213.038	176.212	471.114	127.141	113.252	145.813	82. 555	
	HEMBB1002075				00 708				84. 929
		42.631	31.316	161.071	28.782	21. 239	25. 996	13.087	18. 589
	HEMB81002079	16.958	10. 592	15. 974	7.658	4.913	11.054	12. 406	9. 170
	HEM8B1002080	43.775	32.579	72.576	24.001	9. 827	28.608	17, 214	17.433
	HEMBB1002082	26.775	8. 257	21.193	4. 448	6. 280	19.090	464.903	
20	HEMBB1002084								8. 346
		17, 127	6.840	43. 925	4. 043	9.757	26.316	9. 627	6.512
	HEMB81002088	90.318	38. 977	65.816	40.755	47.974	81.367	57. 452	75. 281
	HEMB81002092	192, 949	59. 522	268. 965	49.978	47.797	60.595	48. 524	38.080
	HEMBB1002094	127.875	84.707	379.671	89.066	80.779	70.636	38, 807	57.037
	HEMB81002103	29.830	9. 307						
				18.867	12.419	117.011	11.825	10. 555	6.133
	HEMBB1002109	28, 380	23. 579	104. 568	24. 307	17.018	17.089	11.301	21.844
25	HEMBB1002115	71.073	86.440	117.523	95. 976	28. 307	85.908	60. 445	114.378
	HEMBB1002120	16.393	10.090	4. 147	2.085	3. 568	9. 594	4, 954	4. 539
	HEMB81002121	12.050	2.757	6. 522					4. 333
					1.146	2.007	0.000	1.999	1.549
	HEMB81002134	784. 781	365. 377	605. 805	262.168	223. 204	719.592	534. 370	450.949
	HEMBB1002136	109, 220	32.405	75.010	27.402	26.278	36. 231	38. 283	23. 593
	HEMBB1002138	17.812	14.057	17. 210	7.413	9. 287	10.613	20.319	9.644
30	HEMBB1002139	51.267	37.549	168.617	27.467	17. 855			
00							27.091	16.428	23. 177
	HEMB81002141	82.369	29. 424	54. 387	14.566	15. 214	39.768	33. 139	22.856
	HEMBB1002142	70.553	42. 309	156. 252	36,636	14.797	26.769	15. 277	22.894
	HEMB81002145	40.661	16. 263	15.725	8.229	13.984	21.757	14.873	15. 525
	HEM881002152	46.728	36.893	105. 608	65.422	40.064	25. 225	29.211	
	HEMB81002162								42. 935
		40.153	34.008	96. 274	29.709	19.847	47.860	22.055	40.550
35	HEMBB1002173	53, 191	41. 151	147.055	26.912	34. 538	16.431	19.449	25. 327
	HEMBB1002189	73.400	88.057	211. 287	73.810	54.029	46.682	45.749	55. 885
	HEMBB1002190	33. 242	51.561	233. 972	49.809	19.665	27. 376	13. 129	61.389
	HEMBB1002193	69, 174	22. 324	33.672	10.803				
						18. 423	27.938	24.748	16. 109
	HEM8B1002217	50. 175	37.602	98.092	38.769	24. 723	33.043	18. 735	39. 436
	HEMBB1002218	596. 902	272.867	712.867	191.461	186. 314	373.711	195. 571	197.556
40	HEMBB1002228	88. 583	45.763	205. 932	47.852	46.693	41.923	37. 485	53.876
	HEMB81002232	56.752	32.790	128.643	36.535	28. 693	32.710	31.447	41.940
	HEMBB1002245	31.084	9.332	17.943	11.049				
						11.834	11.864	17.012	14. 199
	HEMBB1002247	151.502	27.325	64. 167	10.018	26. 829	62.501	35.734	21.698
	HEMBB1002249	153. 327	94.814	380. 989	101.573	65. 579	80.049	62.653	85, 673
	HEMBB1002254	43, 885	36, 756	118, 582	29.328	19.323	11.675	12.693	22. 229
	HEMBB1002255	8.633	2.293	14, 174	8.771	1.813	2. 385	3. 358	
45									3. 589
	HEMBB1002266	5. 303	5.716	8. 530	6.222	1.842	2.404	4.411	2. 295
	HEMBB1002271	160, 682	46.654	157. 828	58. 291	63.843	72.913	62.659	73.702
	HEMBB1002280	24, 597	13.246	76.763	13.976	7.742	9.196	9. 200	16.479
	HEMB81002296	67.004	21.270	52. 536	34. 388	49. 938	53.045	123.030	
	HEMBB1002300		20 600				77.043		41.218
		94.815	28.682	50. 102	35.939	13.923	29.792	25. 246	21.629
50	HEMB81002302	51.059	31.157	28. 441	17.568	17.905	26.026	22.516	30. 501
50	HEM8B1002306	35. 213	49.812	33.017	23, 300	15.072	17. 296	14. 490	16. 293
	HEMBB1002316	19.773	8.638	19.354	3.667	9. 274			
							9, 974	8.613	6.883
	HEMBB1002326	201.896	126.797	406. 052	154.628	89.356	85. 970	54.052	98. 198
	HEMBB1002327	85. 792	48.221	184. 126	47.724	32.764	29. 959	17.415	34. 542
	HEMB81002329	69, 191	21.714	43.746	25.618	17.775	24.892	32.481	27.906
	HEMBB1002340	18.233	28.452	7.730	3.702	3.055			
55		10.200	1 20. 702	1 1.130	3.102	3.055	4. 522	2.914	5.745

Table 48

	HEMB81002342	74.746	83. 579	169.482	40.919	23.495	26. 453	33.215	66. 420
	HEMB81002358	149. 857	132.962	286.214	85.160	50.855	67.646	36.624	78. 432
	HEMBB1002359	160.804	77. 260	219.199	68.995	44. 093	58.049	35. 955	51. 139
5	HEMBB1002364	102.885	74. 409	188.270	50.973	55. 276	45.770	40.780	59.739
	HEMBB1002356	152.074	77.016	248. 465	88.268	81.100	64.637	39. 912	60. 303
	HEMB81002371	44, 433	12. 342	26. 565	13. 307	36.600	10, 553	9. 238	5. 351
	HEMB81002381	134. 427	77. 953	207.310	57.210	48. 215	64. 049	51. 493	77. 629
	HEMB81002383	164, 205	52. 312	94.064	31.346	31.368	30. 947	43. 038	47. 540
	HEMBB1002387	196.859	164. 904	235. 139	49. 485	25. 102	93.004	52. 536	43. 092
10	HEMBB1002409	82.986	49. 978	112.097	29. 207	15. 402	37.667	36.064	38. 132
	HEMBB1002413	123.367	87.690	361.106	87.505	57.485	48. 097	23. 254	49. 302
	HEMBB1002415	87.091	31.703	92. 595	31.804	23. 352	27. 293	21.815	24. 444
	HEMBB1002424	13.162	19. 511	15. 995	5.848	21.533	16.980	18. 246	25. 253
	HEMBB1002425	84.086	69.689	238.147	82.198	36.928	41.171	26.823	47. 957
	HEMBB1002427	143.727	26.894	- 50, 430	25.865	40.707	52. 937	38. 610	47. 517
15	HEMBB1002442	163.853	121. 153	501, 168	129.909	73.231	81.033	47.108	287. 238
	HEMBB1002447	107.214	80.007	214.338	58. 963	41.313	60.452	49. 159	44. 523
	HEMBB1002453	163.250	93.442	384. 443	93.027	68.808	58. 565	46. 254	58. 810
	HEMB81002457	116.756	104.520	330.657	83.026	46.720	50. 971	38. 415	57. 991
	HEMB81002458	18.721	11. 278	23. 232	9.587	7. 205	6.051	4.659	4. 343
	HEMBB1002463	229.657	146.001	663.683	193.622	138.458	104. 827	52.827	110.558
20	HEMBB1002465	44. 210	23. 316	33.631	20.895	17. 932	26.471	19. 122	19.703
	HEMBB1002477	98. 948	27.813	153.875	11.062	36,071	16.072	13.791	8. 347
	HEMBB1002479	23. 249	59.003	73. 224	14.014	10.084	13. 246	1.980	8. 949
	HEMBB1002489	78.748	24.690	71.038	31.400	39.869	43.673	44. 800	75. 957
	HEMB81002492	9.080	6.989	26.130	3. 092	1. 453	5. 606	1.415	2. 381
	HEMBB1002495	95.752	104. 949	301.328	60.728	72.404	45. 161	24. 771	61. 121
25	HEMBB1002502	17. 132	17.866	14.643	16. 170	15. 224	14.056	4. 504	23. 313
	HEMB81002509	0.913	2. 235	7.269	4. 304	0.743	1.283	1, 504	6. 154
	HEMBB1002510	0.732	0.000	0.000	1.858	0. 926	0.000	0.000	0.000
	HEMB81002520 HEMB81002522	249.875	127.604	585. 470 12. 342	169. 423	138. 712 17. 452	90. 360 5. 861	100. 598 8. 292	112.828
	HEMBB1002527	24.741 63.012	27.480 61.066	87.388	46. 392	29. 555	37. 187	25.642	8. 541 36. 089
30	HEMB81002530	72.655	45. 682	83.329	21.750	21.479	53. 227	440. 333	38.710
50	HEMB81002531	40.398	18.832	10.308	9. 953	5. 539	16.743	11.880	8.115
	HEMB81002534	78.552	49. 139	154.741	66.211	30.154	46. 591	28.712	37.112
	HEMB81002536	27.609	22.843	52.264	17.646	8.234	13.078	23. 458	15. 919
	HEMB81002544	24.012	6.185	27.814	13.117	39. 363	15. 921	9. 427	14.017
	HEMB81002545	108.234	31.929	243.949	50.972	16.032	40. 343	31.828	13.472
35	HEMB81002550	31.850	11.452	10.668	11.228	11.049	10, 100	14. 262	14.910
	HEMB81002556	125. 621	89.607	311.607	79.974	50. 209	57.837	53, 696	54.119
	HEMB81002571	33.047	21.526	54. 457	14.847	25. 892	21.961	5. 482	18.608
	HEM881002579	75. 252	55.132	229.479	48.891	31,521	43. 266	24.667	31.554
	HEMB81002582	100, 572	56.574	258.453	63.093	45.740	39, 580	26.474	45. 912
	HEMBB1002584	8. 325	7.614	13.574	6.883	1.796	7.655	6.183	4. 955
40	HEM881002587 HEM881002590	57. 430	44. 383 78. 587	60.900 179.926	47. 981	30.048	30. 562	19.161 33.101	20.854
	HEMBB1002596	114. 241 278. 617	90.944	275.018	65.737	28. 629 68. 247	43.657 114.505	88. 149	34. 032 59. 750
	HEMBB1002600	17.618	16,003	23.907	4. 699	9.726	10, 133	7. 945	8. 940
	HEMBB1002601	67, 910	48. 188	183.948	45. 346	38.021	37. 423	21.860	33, 698
	HEMBB1002603	69. 793	43. 222	141.343	36.733	28.849	35. 264	22.033	29. 436
	HEMBB1002607	64. 941	36. 284	134.598	39. 424	22.220	31.501	15.575	31.024
45	HEMB81002610	22.852	9. 200	51.294	16.832	6.664	12.856	6.433	6.515
	HEMB81002613	85. 026	60.872	161.891	47.532	36.559	44.841	24.569	31.062
	HEM881002614	65.074	30.721	39.687	10,970	15. 910	13. 297	10.461	5. 438
	HEMB81002615	230. 370	55. 581	35.517	11.758	7. 258	46.064	22.857	86.789
	HEMBB1002617	69.016	67.288	254.296	42.530	30.217	36. 395	21.284	37.688
50	HEMB81002623	92.506	78124	204.116	60.739	20.110	48. 078	32.253	43. 355
50	HEMBB1002624	77.755	27.026	163.976	33.209	25.309	20. 104	21.741	24. 486
	HEM881002631	10. 297	18.892	12.879	14.916	7.219	5. 864	6. 990	11.537
	HEMB81002635	88.049	68, 172	141.149	41.853	40.290	23.649	21.781	44. 425
	HEMB81002644	98. 956	65. 380	26.659	19. 268	9. 200	38.890	35.668	29. 597
	HEMBB1002854	127. 571	78.659	51.653	28.747	32.125	137.732	315.048	39.477
55	HEMBB1002661	106. 501	46.651	47.116	19. 470	20.684	30. 561	24. 281	118.028
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Table 49

F	EM8B1002663	100.783	42,600	100.008	36.841	24. 382	35.028	41.975	18.150
	EMBB1002664	179.828	131.008	395.057	51, 235	74, 731	89.130	40.814	55. 262
L.					4, 981	2.314	4. 033	3. 301	1.422
شا ا	EM881002677	2. 206	3.466	5. 138					
{}	EMBB1002683	118. 247	69. 327	247. 117	55. 886	44. 381	26. 944	27.017	42.278
11	EMBB1002684	40. 291	21.056	46.317	17.772	9,039	5.460	8. 120	14.377
	EMBB1002686	30.893	12.882	26.031	19,059	3, 146	12.807	18, 055	9. 131
					29, 779	19.960	25. 893	38.755	15. 268
	EMBB1002692	48, 969	24. 335	52. 440					
jł	EMB81002693	129.760	76.886	322.740	70.620	62.314	67.760	73. 429	39.005
lī l	EM881002697	41.673	38, 793	25. 105	8.999	2.058	7.613	10.266	29.797
	HEMBB1002699	223.756	165.884		116.529	77.378	109.419	79.393	99. 532
				24. 367	3. 561	6. 434	15.699	13. 253	24. 914
	HEMBB1002702	13.506	15.782						
11	HEMB81002705	29. 934	20. 276	16.478	21.230	7. 599	11.487	18.202	30. 589
lī	HEMBB1002712	29.588	10.805	47.572	15.673	13.434	15.691	7. 559	16. 536
	IMR321000028	77.081	39.937	40.934	18.725	8. 281	41.195	27.733	21.472
	IMR321000031	50.644	21.357	34.754	22, 184	15. 786	31.242	22.705	14, 148
L.								26. 605	67. 455
	IMR321000034	76.518	63.230	37. 290	51.243	23.808	43.858		
ſ	IMR321000039	66.895	68.027	83. 136	36, 653	27. 339	62. 232	57. 760	88. 100
Ţ	IMR321000044	1.614	0.000	0.000	0.000	0.000	0.000	1.970	0.000
	IMR321000063	131.633	84.8221	66.499	84.753	43. 262	73.363	69.831	80.878
		157.704	34. 180	42.747	11.752	50.766	66. 106	54. 160	47. 424
	IMR321000085								
į	IMR321000089	52.645	22.980	31.408	17. 365	13.731	36. 296	27. 222	10. 181
ſ	IMR321000091	39.993	32.664	43.895	41.311	25. 143	35. 002	20. 444	63. 906
	LIVER1000004	45.674	30.112	69.445	15.874	11.073	28. 505	106.044	24.660
	LIVER1000008	23.703	14.444	22.304	9. 381	15.657	274.776	344. 333	11. 282
	LIVER1000011	107.957	31. 187	106.032	30. 434	41.030	41.256	348, 474	63, 939
			177.843	270. 232	82. 143	125. 292	206.780	141.934	124. 260
	LIVER1000022	402.839							
	LIVER1000025	61.584	42.776	172.307	36. 300	26.856	33.045	34.820	42. 189
	LIVER1000030	62.987	24.034	69.275	29.784	17. 581	22. 393	51.178	22.556
ì	L1VER1000045	27.941	4.859	27.468	7.384	9.755	14.426	20. 551	24. 802
	LIVER1000046	180. 297	117.998	24.240	23.527	16. 373	7.466	27.795	66.724
•	LIVER1000072	24.097	35. 964	6. 976	11.158	7.657	8.260	16.555	4.898
					13. 193	25. 835	52.139	348.056	37.506
	LIVER1000077	90.518	39. 165	17.306					
	LTVER1000080	17.084	4.918	5. 980	9, 600	2. 294	5. 176	6. 495	4.479
	LIVER1000086	82.711	55. 169	150.708	18.858	19. 278	176.018	481.085	27.747
	LIVER1000092	61.883	36.836	116.592	27.330	16.805	25. 266	35. 863	24. 150
	LIVER1000095	54. 562	13.959	104.146	23.878	13.158	200.163	137.395	5. 508
	LIVER1000097	138. 286	11.401	12.265	8. 127	9.389	9.669	32, 751	7.159
					18. 991	19.124	20.338	142.508	19. 104
	LIVER1000098	58.055	39. 291	47.410					
	LIVER1000100	81.693	64. 546	94. 504	29. 185	18. 588	42.254	23.727	58.633
	LIVER1000101	52. 507	16.303	57.500	10. 286	8.662	17.642	6. 129	27.273
	LIVER1000106	46. 259	32, 121	32.438	11.568	9.377	13.216	102.126	16.904
'	LIVER1000108	26. 277	50, 565	62.172	25. 422	16.619	17.243	38.369	18.508
	LIVER1000115	23. 571	18.673	71.357	14. 244	11.023	17.910	427.626	11.136
	LIVER1000113	100. 902	21.640	35. 183	16. 565	26.236	39.037	87.151	16.249
			27.584	56.479	22.794	25.076	42.015	35, 937	23.833
	LIVER1000138	69.624							
	LIVER1000146	107.757	63.296	209.735	54. 534	42. 231	45. 210	254. 168	42.466
	LIVER1000148	141.467	42. 327	108.510	37, 031	31.920	62. 584	125. 465	65.728
	LIVER1000157	97. 282	37. 198	50.979	49. 952	35.021	43. 954	52.527	43.221
	LIVER1000161	100. 902	24, 883	57.647	28, 329	31.562	42.781	89, 198	30.740
	LIVER1000167	97. 214	29.093	41, 450	25, 700	26.316	112.706	332.789	30.702
	LIVER1000174	53. 927		26.353	13. 595	12.625	36.580	71.460	10.512
							16.773	16.676	14.878
	LIVER1000185	49.746	20.428	31.630	13.964	13.391			
	LIVER1000187	38. 332	8. 211	15.200		8.084	9.846	567.808	8. 320
	LIVER1000190	93. 672	29.635	50.518	15.812	18.768	23.709	41.865	11.496
	LIVER1000192	141.875	53, 337	99.330	32. 936	41.210	79.500	128.608	47.907
	MAMMA1000009	99. 036	77.266	234.005		40.612	44.930	25.218	35.909
	MAMMA1000015	40. 458	7. 192	19.901	13.017	12.921	18.315	13.014	8.185
							42.958	38.148	30.172
	MAMMA1000019	62.999	29.927	150.049		36.450			
	MAMMA1000020	58.696	30.055	181.093		38. 572	34. 176	18, 169	20.807
	MAMMA1000024	15.610	5.088	15, 411	7. 263	3, 468	11.562	37.960	9. 224
	MAMMA1000025	53.708	37.358				24.650	18.530	21. 156
	MAMMA1000043	170. 220	108.774	290.077			82.087	70.843	76. 243
					10.416				8.811
	MAMMA 1000045	83.118	48.873				15.440	7.895	
	NAMMA1000046	117.084	44. 858	285.890	66, 458	43.862	36.388	23. 428	22. 376
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Table 50

AMAMA 1000057 170 331 108 479 221.150 100.356 84.331 77.475 42.047 55.847 148AN 1000058 178 585 50.755 153.319 49.232 88.927 42.539 25.536 52.55 AMAMA 1000058 178 587 50.755 153.319 49.232 88.927 42.539 25.536 52.536 AMAMA 1000058 178 587 50.755 153.319 49.232 88.927 42.539 25.536 52.536 AMAMA 1000058 178 587 59.019 40.608 21.955 13.181 8.822 35.347 56.592 AMAMA 1000058 50.189 20.019 40.608 21.955 13.181 8.822 35.347 56.592 AMAMA 1000059 77.338 37.915 167.474 43.988 16.101 26.961 15.531 22.990 AMAMA 1000059 55.344 38.495 18.888 25.505 11.893 44.990 24.784 25.190 AMAMA 1000059 65.544 54.64 52.522 52.269 24.807 55.730 25.787 23.298 AMAMA 1000059 67.585 32.797 91.551 13.689 19.430 26.8892 22.533 16.804 AMAMA 1000106 67.585 32.797 91.551 13.689 19.430 26.8892 22.533 16.804 AMAMA 1000106 37.916 23.228 90.795 22.075 14.445 24.686 16.649 17.589 AMAMA 1000118 03.7916 23.228 90.795 22.075 14.445 24.686 16.649 17.589 AMAMA 1000118 104.188 58.433 53.822 8.833 24.039 42.731 38.062 43.242 AMAMA 1000133 62.435 25.900 33.061 20.713 14.310 34.686 18.642 14.101 AMAMA 1000133 62.435 25.900 33.061 20.713 14.310 34.686 18.642 14.101 AMAMA 1000133 62.435 25.900 33.061 20.713 14.310 34.686 18.642 14.101 AMAMA 1000134 16.567 8.859 41.797 8.850 17.755 87.598 12.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 78.566 47.352 99.799 34.535 22.772 37.601 28.841 28.800 AMAMA 1000139 79.566 30.800 30.800 30.800 30.700 30.700 30.700 30.700 30.700 30.7									
AMAMA 1000097	MAMMA 1000055	65.118	40.884	57.307	29.859	27. 445	33. 405	22.066	23. 563
AMAMA 100006 79, 588 50, 255 153, 319 49, 223 28, 927 42, 539 25, 616 52, 458 AMAMA 100006 718, 327 15 5010 182, 277, 48, 78, 44, 37, 720 61, 342 45, 357 33, 115 AMAMA 100006 718, 327 15 5010 182, 277, 404 87, 542 63, 176 65, 262 14, 256 45, 922 AMAMA 100006 72 77, 338 37, 915 167, 474 43, 838 16, 101 26, 586 15, 531 22, 393 AMAMA 100006 75, 534 38, 495 38, 888 25, 505 11, 833 44, 990 24, 784 25, 160 AMAMA 100007 55, 534 38, 495 38, 888 25, 505 11, 833 44, 990 24, 784 25, 160 AMAMA 100007 67, 585 32, 797 91, 551 31, 689 19, 430 26, 892 22, 333 16, 642 AMAMA 100010 67 57, 585 32, 797 91, 551 31, 689 19, 430 26, 892 22, 333 16, 642 AMAMA 100010 67 37, 916 23, 228 90, 795 22, 075 14, 445 24, 686 16, 649 17, 569 AMAMA 100010 75 78, 533 24, 45, 600 41, 900 22, 445 16, 140 12, 141 14, 178 AMAMA 100010 75 85, 533 24, 52, 500 18, 835 30, 30, 004 12, 188 31, 1709 11, 461 14, 178 AMAMA 100011 75 86, 533 24, 52, 500 18, 83, 500 30, 004 12, 188 31, 1709 11, 461 14, 178 AMAMA 100011 75 86, 534 33 63, 622 88, 813 45, 90, 22, 181 58, 184 27, 148 15, 487 13, 259 AMAMA 100013 77, 68, 553 27, 255 98, 88, 813 45, 90, 22, 181 58, 179 11, 487 11, 569 AMAMA 100013 77, 62, 555 60, 500 31, 061 20, 713 14, 310 34, 586 18, 642 14, 101 AMAMA 100013 77, 62, 555 60, 500 31, 061 20, 713 14, 310 34, 586 18, 642 14, 101 AMAMA 100013 77, 62, 555 60, 500 31, 061 20, 713 14, 310 34, 586 18, 642 14, 101 AMAMA 100013 77, 62, 555 60, 600 31, 061 20, 713 14, 310 34, 586 18, 642 14, 101 AMAMA 100013 77, 62, 555 60, 600 31,	MANMA1000057		108, 479	421, 160	100.366	84, 331	77, 475	42, 047	55, 847
AMAMA1000059 118.391 15.010 182.772 48.764 43.720 61.342 45.357 33.115 AMAMA1000085 40.199 20.019 40.608 21.955 13.181 18.822 36.347 26.2784 AMAMA1000097 77.338 37.915 167.474 43.888 16.101 26.961 15.531 22.390 AMAMA1000097 62.346 34.8495 38.888 25.605 11.893 44.590 24.784 25.180 AMAMA1000097 62.346 34.8495 38.888 25.605 11.893 44.590 27.784 25.180 AMAMA1000097 62.346 34.8495 38.888 25.605 11.893 44.590 27.784 25.180 AMAMA1000097 62.346 34.8495 38.888 25.605 11.893 44.590 27.784 25.180 AMAMA1000098 55.344 38.495 38.888 25.605 11.893 44.590 27.784 25.180 AMAMA1000098 67.585 37.2797 91.55 31.689 19.430 26.892 22.353 16.847 AMAMA1000108 63.752 26.301 88.530 30.004 12.188 31.709 11.461 14.718 AMAMA1000117 58.533 24.502 43.190 22.445 16.140 27.418 15.487 13.269 AMAMA1000117 58.533 24.502 43.190 22.445 16.140 27.418 15.487 13.269 AMAMA1000118 104.188 88.431 53.822 8.833 42.039 42.731 18.062 43.242 AMAMA1000118 104.188 88.431 53.822 8.833 42.039 42.731 18.062 43.242 AMAMA1000118 36.275 25.090 33.001 20.713 14.310 43.666 18.642 14.101 AMAMA1000119 78.555 74.256 98.813 45.970 22.181 58.739 90.197 14.587 AMAMA1000119 78.565 72.256 98.813 45.970 22.181 58.739 90.197 14.587 AMAMA1000119 78.565 78.258 12.8150 13.910 5.510 14.314 12.120 15.748 AMAMA1000118 30.212 20.528 28.150 13.910 5.510 14.314 12.120 15.748 AMAMA1000118 30.121 20.528 28.1860 13.910 5.500 127.758 76.596 45.255 60.360 AMAMA1000118 30.121 20.528 28.1860 13.910 5.500 127.758 76.596 45.255 60.360 AMAMA1000118 30.121 20.528 38.888 57.289 28.89 60.007 66.037 34.872 51.888 AMAMA1000119 30.121 20.528 38.888 57.289 38.89 60.007 66.037 34.872 51.888 AMAMA1000118 30.121 20.528 38.888 57.289 38.89 60.007 66.037 34.872 51.888 38.888 57.289 38.898 60.007 66.037 34.872 51.888 38.888 57.289 38.888 60.007 66.037 34.872 51.888 38.888 57.289 38.888 51.889 60.007 66.037 34.872 51.888 38.888 57.289 38.888 57.289 58.888 50.007 66.037 34.872 51.888 38.888 57.289 57.389 38.888 57.289 57.389 38.888 57.289 57.389 38.888 57.389 58.889 58.889 58.399 58.399 58.3									
AMAMA 1000084 128. 354 92. 319 277. 404 87. 542 63. 176 65. 252 34. 266 45.092 AMAMA 1000085 0. 139 20. 019 40. 608 21.956 13. 181 8. 822 36. 347 72. 52. 390 AMAMA 1000085 55. 344 38. 495 38. 888 25. 605 11. 893 44. 990 24. 784 25. 180 AMAMA 1000087 65. 546 54. 54. 54. 54. 54. 54. 54. 54. 54. 54.									
AMALIODO085									
LAMAL 1000192 77, 338 37, 915 167, 376 43, 988 16, 101 25, 961 15, 551 22, 390 LAMAL 1000095 55, 344 38, 495 38, 888 25, 505 11, 893 44, 990 24, 784 25, 160 LAMAL 10001967 65, 565 54, 54, 54, 54, 54, 54, 54, 54, 54, 54									
AMANI 10000196 55, 344 38, 495 38, 888 25, 605 11, 893 44, 990 24, 784 25, 160 AMANI 10000197 62, 546 54, 694 55, 527 52, 527 52, 699 24, 807 65, 730 57, 787 23, 381 AMANI 1000102 67, 585 32, 797 91, 551 31, 689 19, 430 25, 892 22, 353 16, 842 AMANI 1000103 63, 795 22, 263, 301 88, 835 30, 300, 41, 21, 888 31, 709 11, 461 14, 718 AMANI 1000103 63, 795 22, 228 30, 795 22, 075 14, 445 24, 686 16, 649 17, 589 AMANI 1000117 8, 85, 33 24, 502 43, 190 22, 445 16, 140 27, 418 15, 487 13, 269 AMANI 1000118 104, 188 58, 433 63, 822 8, 833 32, 039 42, 731 38, 1062 43, 224 AMANI 1000118 104, 188 58, 433 63, 822 8, 833 32, 039 42, 731 38, 1062 43, 224 AMANI 1000118 104, 188 52, 50, 900 33, 30, 61 20, 713 14, 310 4, 686 18, 642 14, 104 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 522 79, 900 246, 344 90, 530 127, 758 76, 596 45, 125 60, 360 AMANI 1000118 105, 720 76, 720 77									
AMAMA 1000197 62, 546 54, 654 52, 522 52, 269 24, 807 55, 730 25, 787 23, 298 AMAMA 1000108 67, 555 32, 797 91, 555 31, 869 91, 93, 90 26, 892 22, 353 16, 842 24, 244 24, 246 25, 246 24, 246									
ALMAN ALMON ALMAN ALMA									
AMMATIODOTIOS 63.752 25.301 89.530 30.004 12.188 31.709 11.61 14.718 ALMMATIODOTIOS 37.916 23.228 90.795 22.075 14.456 24.666 16.649 17.569 ALMMATIODOTIOS 58.533 24.502 43.190 22.445 16.140 27.418 15.487 13.289 ALMMATIODOTIOS 19.168 58.433 33.822 8.833 24.039 42.731 33.062 43.242 ALMMATIODOTIOS 170.665 72.256 88.813 45.970 22.181 58.739 50.197 14.587 ALMMATIODOTIOS 170.665 72.256 88.813 45.970 22.181 58.739 50.197 14.587 ALMMATIODOTIOS 19.45 25.909 33.061 20.713 14.310 34.686 18.642 14.101 ALMMATIODOTIOS 19.566 47.362 99.179 34.535 22.772 37.601 28.841 28.280 ALMMATIODOTIOS 19.566 47.362 99.179 34.535 22.777 37.601 28.841 28.280 ALMMATIODOTIOS 16.647 8.669 41.797 8.690 9.949 10.059 4.040 8.280 ALMMATIODOTIOS 20.503 88.642 28.1247 110.848 80.817 97.755 63.045 78.585 ALMMATIODOTIOS 20.503 88.642 28.1247 110.848 80.817 97.755 63.045 78.585 ALMMATIODOTIOS 52.05.33 88.642 28.1247 110.848 80.817 97.755 63.045 78.585 ALMMATIODOTIOS 52.05.03 88.642 28.1247 110.848 80.817 97.755 63.045 78.585 ALMMATIODOTIOS 19.168 8.683 77.390 22.848 21.852 41.672 11.036 10.618 ALMMATIODOTIOS 19.168 8.683 77.390 22.848 21.852 41.672 11.036 10.6181 ALMMATIODOTIOS 19.168 8.683 77.390 22.848 21.852 41.672 11.036 10.6181 ALMMATIODOTIOS 19.316 8.683 77.390 22.848 21.852 41.672 11.036 10.6181 ALMMATIODOTIOS 19.316 8.683 77.390 22.848 21.852 41.672 11.036 10.6181 ALMMATIODOTIOS 19.316 8.683 77.390 22.848 21.852 41.672 11.036 10.6181 ALMMATIODOTIOS 19.316 8.683 77.390 42.850 23.750 23.680 21.050 46.992 ALMMATIODOTIOS 19.316 8.683 77.390 42.850 23.750 23.680 21.050 46.992 ALMMATIODOTIOS 19.316 8.683 77.390 42.850 23.750 23.880 21.050 46.992 ALMMATIODOTIOS 19.316 8.683 77.390 42.850 23.750 23.880 27.779 41.670 27.770 47.780 47.7	MAMMA1000097	62.546	54.694	52. 522	52.269	24.807		25. 787	23. 298
AMBAI 1000103 63.752 26.301 88.530 30.004 12.188 31.709 11.61 14.718 AMBAI 1000107 58.532 24.507.55 14.445 24.566 16.649 17.569 AMBAI 1000117 58.533 24.507 43.190 22.445 16.140 27.418 15.487 13.289 AMBAI 1000118 10.188 38.433 63.822 8.833 24.039 42.7418 15.487 13.289 AMBAI 1000118 10.6522 72.256 98.813 45.970 22.181 58.739 50.197 14.587 AMBAI 1000113 10.6522 79.909 24.6534 90.500 12.713 14.310 34.686 18.642 14.101 14.581 14.5	MAMMA1000102	67.585	32.797	91.551	31.689	19.430	26.892	22. 353	16.842
LAMAN 1000117	MAMMA1000103	63.752	26. 301	89.530	30.004	12.188	31.709	11.461	14.718
ALMAN 1000175 58.533 24.502 43.190 22.445 16.140 27.418 15.487 13.269 ALMAN 1000179 170.655 72.256 38.313 45.970 22.181 58.739 50.197 41.587 ALMAN 1000139 170.655 72.256 38.313 45.970 22.181 58.739 50.197 41.587 ALMAN 1000134 106.522 79.990 246.344 90.550 127.758 76.596 45.225 60.360 ALMAN 1000134 106.522 79.990 246.344 90.550 127.758 76.596 45.225 60.360 ALMAN 1000134 30.121 20.5281 28.150 13.910 5.510 14.314 12.120 15.748 ALMAN 1000135 78.566 47.362 99.179 34.535 22.772 37.601 28.841 28.280 ALMAN 1000143 16.647 8.669 41.797 8.690 9.949 10.059 4.040 8.280 ALMAN 1000155 205.301 38.642 281.247 118.84 80.817 97.755 63.045 78.585 ALMAN 1000155 205.301 38.642 281.247 118.84 80.817 97.755 63.045 78.585 ALMAN 1000165 205.301 38.642 281.247 118.84 80.817 97.755 63.045 78.585 ALMAN 1000175 13.625 68.928 55.7239 22.848 21.852 41.672 11.036 10.618 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.340 52.609 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.340 52.609 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.340 52.609 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.340 52.609 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.340 52.609 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.840 ALMAN 1000175 103.027 21.955 68.080 33.572 25.668 45.271 40.850 ALMAN 1000175 103.605 33.830 36.836 29.409 38.905 35.131 35.059 36.6570 ALMAN 1000175 33.830 36.823 36.836 29.409 38.905 35.131 35.059 36.6570 ALMAN 1000175 33.830 36.823 36.836 29.409 38.905 35.131 35.059 36.6570 ALMAN 1000276 46.455 59.079 71.789 26.771 29.275 55.156 32.232 49.295 ALMAN 1000278 33.	MANSA 1000106	37.916	23, 228	90, 795	22.075	14. 445	24. 686	16.649	17. 569
ALMAN 1000 18 104 168 554 433 53 822 8,833 24 039 42 731 33 062 43 242 ALMAN 1000 133 62 435 72 556 98 813 45 790 22 181 58 739 50 197 14 587 ALMAN 1000 133 62 435 25 590 33 061 20 713 14 310 34 586 18 642 14 101 ALMAN 1000 134 78 566 47 362 99 179 34 535 22 772 37 501 28 341 38 28 ALMAN 1000 139 78 566 47 362 99 179 34 535 22 772 37 501 28 341 38 28 ALMAN 1000 141 30 12 20 5281 28 150 13 910 5 510 14 314 12 120 15 748 ALMAN 1000 134 16 647 68 689 41 797 8 650 9 94 10 059 4 040 8 280 ALMAN 1000 135 20 50 31 88 642 291 247 110 884 80 817 97 75 5 50 64 78 28 ALMAN 1000 155 20 50 31 88 642 291 247 110 884 80 817 97 75 5 50 64 78 58 ALMAN 1000 150 128 128 259 413 21 24 24 28 27 27 86 623 42 257 5 63 04 78 58 ALMAN 1000 151 41 425 45 928 255 746 98 189 60 007 66 037 34 87 25 50 10 ALMAN 1000 175 10 30 20 21 955 68 80 30 37 57 25 68 45 27 40 340 52 609 ALMAN 1000 175 19 316 8 683 7 960 4 530 3 535 7 894 5 5 5 5 60 ALMAN 1000 175 19 316 8 683 7 960 4 530 3 535 7 894 5 5 5 6 6 6 27 ALMAN 1000 191 8 72 21 555 68 80 30 20 20 20 20 20 20 2	MAMMA1000117			43, 190	22.445		27.418	15, 487	
ALMAN 1000125 170, 655 72, 256 98, 813 45, 970 22, 181 58, 739 50, 197 14, 587 ALMAN 1000133 62, 435 25, 909 33, 061 62, 713 14, 310 34, 686 18, 642 14, 101 ALBIN 1000134 106, 522 79, 080 246, 344 90, 530 127, 758 76, 596 45, 325 60, 360 ALBIN 1000135 78, 566 47, 352 99, 179 34, 535 22, 772 37, 601 28, 841 28, 280 ALBIN 1000134 30, 121 20, 5281 28, 150 13, 910 5, 510 14, 314 12, 120 ALBIN 1000135 128, 128 259, 413 21, 844 28, 280 ALBIN 1000143 16, 647 8, 669 41, 797 8, 690 9, 949 10, 059 4, 040 82, 280 ALBIN 1000150 128, 128 259, 413 21, 844 28, 287, 77 86, 623 42, 227 51, 840 ALBIN 1000163 128, 128 259, 413 21, 844 28, 277 86, 623 42, 227 51, 840 ALBIN 1000163 128, 128 259, 413 21, 844 28, 877 86, 623 42, 227 51, 840 ALBIN 1000163 128, 128 259, 413 21, 844 28, 877 86, 623 42, 227 51, 840 ALBIN 1000163 13, 613 36, 838 57, 239 22, 848 21, 852 41, 672 11, 018 10, 618 ALBIN 1000173 141, 225 46, 928 585, 746 98, 189 60, 007 66, 037 34, 875 ALBIN 1000173 19, 316 8, 683 7, 950 45, 592 ALBIN 1000173 19, 316 8, 683 7, 950 43, 592 ALBIN 1000193 57, 490 53, 830 148, 702 42, 892 23, 250 23, 680 21, 050 45, 992 ALBIN 1000191 88, 722 31, 449 40, 834 26, 664 22, 392 26, 766 36, 253 27, 729 ALBIN 1000192 53, 457 25, 036 30, 205 28, 380 21, 976 101, 288 128, 339 44, 025 ALBIN 1000193 30, 326 36, 823 36, 836 29, 409 18, 905 35, 131 35, 059 36, 667 ALBIN 1000194 30, 212 39, 338 36, 823 36, 836 29, 409 18, 905 35, 131 35, 059 36, 667 ALBIN 1000204 64, 455 59, 079 71, 789 26, 771 29, 275 55, 156 67, 220 38, 836 ALBIN 1000204 64, 455 59, 079 71, 789 26, 771 29, 275 55, 156 67, 220 38, 38 ALBIN 1000224 50, 365 20, 174 11, 901 11, 838 17, 236 39, 30 31, 377 ALBIN									
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MANNA1000251 115. 401 47. 888 209. 360 39. 959 42. 597 57. 904 34. 572 51. 015				1					
MAMMA1000251 115.401 47.888 209.360 39.959 42.597 57.904 34.572 51.015 MAMMA1000254 43.161 20.910 114.081 20.548 9.699 9.885 5.346 32.024 MAMMA1000257 142.781 70.118 332.822 104.425 84.387 124.673 78.270 116.103 MAMMA1000262 18.952 34.301 19.786 32.516 14.840 15.513 23.805 35.519 MAMMA1000264 59.532 20.630 124.043 44.847 29.466 21.390 22.616 37.039 MAMMA1000265 55.476 28.959 122.654 35.663 27.018 24.021 20.212 38.284 MAMMA1000270 142.968 64.234 270.948 75.022 64.760 68.130 64.006 73.994 MAMMA1000271 53.605 9.611 35.682 12.139 16.139 24.236 26.722 26.433 MAMMA1000277 56.407 16.435 98.448 19.751 12.725 33.047 23.839 33.012 MAMMA1000279 68.661 36.984 173.379 46.809 34.441 42.500 26.143 48.597 MAMMA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MAMMA1000309 17.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744	MAMMA1000245								
MANNA1000254									
MAMMA1000257	MAMMA1000251								
MANNA1000262 18.952 34.301 19.786 32.516 14.840 15.513 23.805 35.519 MANNA1000264 59.532 20.630 124.043 44.847 29.466 21.390 22.616 37.039 MANNA1000265 55.476 28.959 122.654 35.663 27.018 24.021 20.212 38.284 MANNA1000270 142.968 64.234 270.948 75.022 64.760 68.130 64.006 73.994 MANNA1000271 53.605 9.611 35.682 12.139 16.139 24.236 26.722 26.433 MANNA1000277 56.407 16.435 98.448 19.751 12.725 33.047 23.839 33.012 MANNA1000278 40.286 13.365 19.395 9.730 12.609 20.423 25.204 22.237 MANNA1000279 68.661 36.984 173.379 46.809 34.441 42.500 26.143 48.597 MANNA1000283 55.199 27.095 46.168 22.395 15.870 21.308 16.298 18.504 MANNA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MANNA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MANNA1000284 457.450 361.106 313.407 116.696 112.848 343.951 155.948 100.375 MANNA1000298 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MANNA1000302 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 MANNA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MANNA1000309 10.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744	MAMMA1000254								
MAMMA1000264 59.532 20.630 124.043 44.847 29.466 21.390 22.616 37.039 MAMMA1000265 55.476 28.959 122.654 35.663 27.018 24.021 20.212 38.284 MAMMA1000271 53.605 9.611 35.682 12.139 16.139 24.236 26.722 26.433 MAMMA1000277 53.605 9.611 35.682 12.139 16.139 24.236 26.722 26.433 MAMMA1000277 56.407 16.435 98.448 19.751 12.725 33.047 23.839 33.012 MAMMA1000278 40.286 13.365 19.395 9.730 12.609 20.423 25.204 22.237 MAMMA1000278 40.286 13.365 19.395 9.730 12.609 20.423 25.204 22.237 MAMMA1000279 68.661 36.984 173.379 46.809 34.441 42.500 26.143 48.597 MAMMA1000283 55.199 27.095 46.168 22.395 15.870 21.308 16.298 18.504 MAMMA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000288 457.450 361.106 313.407 16.696 112.848 343.951 155.948 100.375 MAMMA1000298 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MAMMA10003002 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 MAMMA10003007 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MAMMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744 MAMMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744 MAMMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744 MAMMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744 MAMMA1000309 279.600 275.000	MAMMA1000257							78.270	
MANIMA1000270	MAMMA1000262								
MAIMA1000270	MAMMA 1000264								
MAIMA1000271 53.605 9.611 35.682 12.139 16.139 24.236 26.722 26.433 MAMMA1000277 56.407 16.435 98.448 19.751 12.725 33.047 23.839 33.012 MAMMA1000279 40.286 13.365 19.395 9.730 12.609 20.423 25.204 22.237 MAMMA1000279 68.661 36.984 173.379 46.809 34.441 42.500 26.143 48.597 MAMMA1000283 55.199 27.095 46.168 22.395 15.870 21.308 16.298 18.504 MAMMA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MAMMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAMMA1000284 457.450 361.106 313.407 116.696 112.848 343.951 155.948 100.375 MAMMA1000280 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MAMMA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MAMMA1000303 57.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MAMMA1000305 32.363 19.693 108.733 15.375 12.695 14.455 13.353 15.189 MAMMA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MAMMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744	MAMMA1000266								
MAIMIA1000277 \$6, 407 16, 435 98, 448 19, 751 12, 725 33, 047 23, 839 33, 012					75.022				
MAMMA1000278	MAMMA1000271								26. 433
MANNA1000279 68.661 36.984 173.379 46.809 34.441 42.500 26.143 48.597 MANNA1000283 55.199 27.095 46.168 22.395 15.870 21.308 16.298 18.504 MANNA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MANNA1000284 457.450 361.106 313.407 116.696 112.848 343.951 155.948 100.375 MANNA1000298 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MANNA1000302 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 MANNA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MANNA1000305 32.363 19.693 108.733 15.375 12.695 14.455 13.353 15.189 MANNA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MANNA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744	MAMMA1000277	56, 407			19.751				33.012
MAIMA1000283 55.199 27.095 46.168 22.395 15.870 21.308 16.298 18.504 MAIMA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MAIMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAIMA1000287 457.450 361.106 313.407 116.696 112.848 343.951 155.948 100.375 MAIMA1000298 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MAIMA10003002 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 40.441	MAMMA1000278								
MAIMA1000284 76.726 67.676 42.784 39.851 34.586 47.651 39.169 48.342 MAIMA1000287 73.583 58.726 142.953 39.301 31.007 27.370 29.006 35.599 MAIMA1000294 457.450 361.106 313.407 116.696 112.848 343.951 155.948 100.375 MAIMA1000298 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MAIMA1000302 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 40.86	MAMMA1000279								
MAIMA1000287	MAMMA1000283		27.095		22. 395		21.308		18. 504
MAMMA1000294 457.450 361.106 313.407 116.696 112.848 343.951 155.948 100.375	MAMMA1000284	76.726	67.675						
MAIMIA1000294	MAMMA1000287			142.953		31.007			35. 599
MAMMA1000298 31.731 25.511 41.413 16.220 16.320 14.676 22.043 20.205 MAMMA1000302 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 MAMMA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MAMMA1000305 32.363 19.693 108.733 15.375 12.695 14.455 13.353 15.189 MAMMA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MAMMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744	MAMMA1000294			313.407	116.696	112.848			
MANNA1000302 109.379 58.532 280.880 69.156 44.790 36.788 28.220 40.861 MANNA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MANNA1000305 32.363 19.693 108.733 15.375 12.695 14.455 13.353 15.189 MANNA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MANNA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744	MAMMA1000298				16, 220	16. 320	14.676		
MANNA1000303 67.505 14.147 18.804 11.073 33.859 26.599 30.177 30.810 MANNA1000305 32.363 19.693 108.733 15.375 12.695 14.455 13.353 15.189 MANNA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MANNA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744									
MANNA1000305 32.363 19.693 108.733 15.375 12.695 14.455 13.353 15.189 MANNA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MANNA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744									
MANMA1000307 279.600 75.098 397.421 75.020 45.244 68.757 131.117 116.800 MANMA1000309 11.679 39.455 13.529 3.502 3.904 8.895 10.500 6.744									
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		1 22.043				1 01 100		,	

Table 51

	MAMMA 1000313	79. 577	69.550	54, 317	10.741	60.526	42.964	18. 206	37. 303
	MAMMA1000331	80.910	48.868	139.047	33.811	22.564	·15. 207	18. 580	21. 385
	MAMMA1000335	54. 800	22.399	33.190	18. 244	16.273	30. 688	26.611	30.790
	MAMMA1000339	69. 222	40.948	83.679	13. 158	20.941	22. 134	20. 026	10.739
	MAMMA1000340	57. 498	34.708	164.968	32, 922	28.610	23.069	18. 858	23.519
	MAMMA1000348					32, 546			
		78.099	102.955	374.737	55.033		66. 256	22. 303	23. 575
	MAMMA1000356	152. 238	116.086	454.516	67. 232	34. 525	47.884	22. 865	61. 267
	MAMMA 1000358	34, 367	56.332	15, 362	15.091	15.743	17. 405	19.645	7. 358
	MAMMA 1000360	71.104	74.351	246.244	43.414	24.093	24. 945	14. 842	14. 739
t .									
	MAMMA 1000361	101.653	93.468	230. 215	73.577	45.022	37. 236	37. 987	42. 992
	MAMMA 1000363	71.108	19. 232	39.013	13.717	23.713	30.73 9	27.813	32, 485
	MAMMA1000370	171.867	108.830	110.466	80.949	52.076	79.266	57.877	247.810
	MAMMA1000371	100. 543	32.223	80.873	48.039	49.442	91.739	57. 647	
									46. 599
	MANNA 1000372	206.850	114. 326	609.068	130. 138	79.980	80.890	54. 857	97.509
	MAMMA 1000385	72.074	60, 911	238. 462	40.061	34. 528	31.361	22. 458	45. 681
	MAMMA1000388	118.855	69.094	105.789	42.626	50.059	55. 389	37. 396	37.825
	MAMMA1000395	97.031	44. 493	34. 493	20. 201	19,036	27. 695	24. 269	
									17. 433
	MAMMA 1000402	126. 085	107.637	256. 584	68.415	45.669	61.486	<u>30.</u> 340	30. 943
	MAMMA1000403	87. 558	63.749 (208. 574	64.857	45. 578	44. 799	22.710	42. 239
	MAMMA1000410	43.073	43.539	94. 207	39.613	19.880	22.573	16. 272	21.003
	MAMMA1000413	30.829	13. 370	70.418	17. 102	13.392	15. 291	11. 599	
ı									15. 353
	MAMMA1000414	125. 550	111.622	81.672	15.722	51.528	14. 549	28. 214	13.858
	MAMMA1000416	179.864	103.793	427.214	107.383	105.899	121. 441	55.040	84.667
	MAMMA1000421	131.712	73.475	307.780	70.841	55.037	49.498	34. 519	46, 482
	MAMMA1000422	12.614	14. 628	30.167	16.100	11.675	22.441	18.843	54. 831
	MAMMA1000423	34.100	22. 150	69.677	18.461	13.815	15.645	8. 500	8.869
	MAMMA1000424	9. 330	4. 056	36.234	8. 171	0.971	2.769	0.745	7.267
	MAMMA1000429	575. 321	219.603	317.414	158. 529	150,779	290. 300	196. 161	149.619
	MAMMA1000431	143. 825	79.993	275.497	82.499	52.496	63. 425	43.337	66.733
	MAMMA1000432								
		65. 212	17.117	24.472	28.083	17.360	33. 881	27.547	29.615
	MAMMA1000437	89. 375	88. 947	265. 572	60.025	69.885	45. 195	30.823	31.510
	MANMA 1000444	120.017	124, 234	477.772	115.966	65. 200	66.888	31.943	88.274
	MAMMA1000446	50. 201	66.027	41.406	8.991	18.971	29. 395	7.985	37. 220
	MAMMA1000449								
		81.386	41. 427	180.761	40.414	25, 983	35. 232	23. 109	27. 942
	MAMMA1000457	47.862	13.862	15.095	11.981	7.566	21.142	12.971	10.872
	MAMMA1000458	34. 485	13.749	22.864	12.116	11.199	18.881	15. 924	10.046
	MAMMA1000468	8. 235	7.843	6.029	5.004	5. 503	8. 258	7.138	1.618
	MAMMA1000472	250. 243	67.964	110.774	68.614	73.186	111.758	88.016	
									79. 409
	MAMMA1000473	54. 174	16.506	40. 489	16.002	17.450	26.506	17.741	13.900
1	MAMMA1000477	77.316	50.237	238.943	56.460	38.807	32.776	36.438	35.332
	MAMMA1000478	201.299	157.097	496.514	127.872	82.832	77.444	49. 296	86.763
	MAMMA1000483	107.340	74.564	252.463	60.824	31.055	44. 198	44. 167	87. 449
	MAMMA 1000490								
		14.473	14.068	16.023	12.496	8.202	15.654	11.091	12. 344
	MAMMA 1000496	32.756	10.554	20.693	10.676	19.830	19. 282	13. 204	13.410
	MAMMA 1000500	23.016	17.584	49.151	15.706	13.914	19.063	11.094	22.904
	MAMMA1000501	196.637	102.490	468.793	104. 118	67.761	83.834	76.446	86.912
	MAMMA1000503	7.083	4.085	3.865	1.004	1.005	3.752	4.005	3. 248
	MAMMA1000506	201.452	116.279	151, 434	56.847	78.502			
							149.780	99. 352	64.069
	MAMMA1000510	70.898	18.432	60.927	39. 187	33. 327	42.829	40. 993	33. 127
	MAMMA1000515	43.923	30.031	85.637	35.744	18.805	21.837	19. 339	17.922
	MAMMA1000516	74.742	48.811	148. 307	43.452	18.069	34.061	19. 122	26. 985
	MAMMA1000522	53. 273	23.845	132.197	22.861		24.776		
:		33.273				14.594		12.095	27.578
!	MAMMA1000524	130.806	61.389	266. 529	71.558	50.972	73.691	47. 484	55. 510
	MAMMA1000528	38. 579	27.136	46.940	35.839	15.860	29.316	19.300	24.797
	MAHMA1000534	32.603	20.088	33.950	10.973	7.185	10.580	7.972	10.160
	MAMMA1000541	165. 518	58.806			27.705	52, 036	46. 200	39.018
				85.648	63. 188				
	MAMMA 1000550	119.597	203.059	41, 184	24. 393	5.859	48. 433	766.194	63.005
	MAMMA 1000556	31.963	15.056	15. 588	8.634	11.294	15. 698	21.467	16.597
)	MAMMA 1000559	57.738	31.181	242.155	29.443	19,030	26.908	13.520	41.571
	MAMMA 1000565	118.770	30.318	289. 829	37.509	33.728	38.720	18. 344	26.847
	MAMMA1000567	77.050	44. 379	224.645	48. 804	41.102	56.039	36. 496	63. 529
	MAMMA 1000576	271.038	180.600	661.566	221.987	157. 443	132. 385	93.679	129.843
	MAMMA 1000582	54, 936	43.406	272. 366	14. 342	18.896	29. 396	46.333	40.210
	MAMMA 1000583	90. 692	51.670	147. 946	34. 905	17. 175	23. 177	19.077	40.824
		30.032	71.010	141.340	1 34. 303	1 11.113	69.111	13.011	40.024

Table 52

			*** *** 1	- CO 0CO 1	20 242	30 100 T	24 000 1	46.734
MANGKA1000585	89.865		288. 673	52. 259	29. 243	39. 188	24.088	
MAMMA1000587	47.955	14. 789	58. 279	12.415	6. 584	14.410	15.734	6.826
MAMMA1000591	77.705	38. 280	81.784	28.019	20.094	28. 578	24. 299	19.949
MAMMA1000594	194. 593	94. 384	488. 898	91.064	59. 244	55.681	43.577	75.029
MAMMA1000597	496.923	264. 906		196. 294	121.483	306.397	199.968	160.426
MAMMA1000605	324. 584	183.667		209. 555	135.844	158.096	97.598	149. 183
MAMMA1000612	68.113	22.051	42. 999	14.074	19. 294	41.220	29.460	15.713
MAMMA1000614	580.099	135.874	402.890	69.022	127.808	309.892	249.344	194.110
MAMMA1000616	2.590	16.442	13.809	1, 109	3.011	7.500	3.036	3.188
MAMMA1000621	19. 258	12.723	14. 307	13.200	5. 971	12.028	11.561	11.081
MAMMA1000623	50.189	23. 285	25. 913	12.057	10.648	23. 327	19.218	20.667
MAMMA1000625	651.334	249. 117	346.876	155. 944	192.671	373.924	300. 473	274.263
MAMMA1000625	4. 459	2.994	4. 756	2.883	0.000	4.118	5. 584	9. 542
MAMMA1000643	24, 259	51.698	115. 511	47.881	17.554	52.330	16.308	38.448
MAMMA1000645	72.487	111, 121	22.868	9. 213	27.074	81.604	46.859	34.048
MAMMA1000652	152. 920	94. 568	319. 943	76.610	67.817	87.605	41.747	77.720
		41.097	278.504	38. 131	36. 289	67.327	34. 224	32. 593
MAMMA 1 000657	116.830	37.993	133.863	26.712	16.308	21. 135	14, 102	35. 215
MAMMA1000664	48.908			25. 027	29. 493	43.769	22.193	24. 502
MAMMA1000667	77.285	24.312	99.732	17. 454	18. 336	50, 398	38. 233	26. 553
MAMMA1000668	42.561	28.100	54. 970			12.841	7.392	12. 088
MAMMA1000669	22.797	14.382	57. 803	14.670	6.337 25.826	33.332	38.768	39.130
MAMMA1000670	66.748	22.566	46.836	26. 498			38, 934	40. 592
MAMMA1000672	128. 331	25. 209	67.913	35. 262	28.783	64.713		
MAMMA1000681	66.397	40,677	32.249	14.404	13. 181	26.710	30.054	37. 369
MAMONA 1000684	85. 908	107.381	66.100	35. 992	32.881	41.006	36.719	77. 834
MAMMA 1000696	165, 293	107.442	551.458	130.714	88.510	70.985	43.857	55. 551
MAMMA1000702	82.316	25.689	52.797	22.639	22.884	48.899	39.297	29.636
MAMMA1000706	81.416	25. 442	34. 529	20. 432	15, 562	39.909	33.303	25. 371
MAMMA 1000707	128. 277	17, 100	51.835	15.001	33.473	48.628	46. 555	24. 075
MAMMA 1000713	75.263	59.677	109.995	37. 970	23. 975	33.874	30.149	39. 491
MAMMA1000714	228.366	288.017	246.261	56.045	25. 380	80.480	51.219	64. 589
MAMMA1000718	98, 208	92.149	245.750	79. 940	49.064	50. 180	40.223	49. 032
MAMMA 1000720	158.737	111.227	446.586	101.175	73.612	78.021	29.904	60. 252
MAMMA1000723	64.930	49.053	148. 286	40. 276	28.806	19.434	18.845	24. 784
MANMA 1000731	31.516	11.357	68.834	12. 436	11.755	7.989	7.536	7. 36
MAMMA1000732	121. 291	56.513	230.064	68.746	51.582	53.763	35. 440	49. 33
MAMMA1000733	24. 525	14. 171	58.717	16, 852	7.153	14.100	8. 586	10.63
MAMMA1000734	113.011	127.466	142.152	102.345	44.860	84.456	43.098	98.01
MAMMA1000736	142.978	48.490	130.520	34. 595	40.252	73.418	82.810	69. 46
MAMMA1000738	110.304	61.504	28.831	38. 642	18.942	31.735	48. 926	35. 12
MAMMA 1000744	140.264	94. 669	281. 287	76. 261	79.000	63.977	43. 557	40. 38
MAMMA1000746	26.385	50.110	37.264	16.895	10.790	35. 280	3. 177	11.01
MAMMA 1000748	73.879	36.619	52.587	30. 957	36.810	46.899	25. 359	24.84
MAMMA1000751	42.505	27.882	58.087	44. 924	28. 537	43.075	32. 581	61.05
MANMA1000752	55. 785	55.799	193.100	53.436	25.798	29.655	21.969	44. 38
MAMMA1000757	314.709	210.647	536.246	187.416	161. 327	151.926	112.625	152.07
MAMMA1000760	218.937	178.377	534. 346	131.736	100.173	95. 443	58. 158	91.22
MAMMA1000761	147, 993	73.793	349.399	85. 319	65.436	75. 180	43.310	63.42
MAMMA1000775	75.873	25.684	170,040	34. 150	30.063	20.938	15.825	18. 99
MAMMA 1000776	101.206	81.986	253.211	57.436	51.043	51.597	28. 394	33.45
MAMMA1000778	71.839	47.596	214, 100	42.749	28.124	29.701	17.866	26.49
MAMMA 1000781	67.901	30.437	97.580		23. 265		17. 488	26.97
MAMMA 1000782	286.062	65.796	174.951	84.753			90.446	85.36
MAMMA1000784	135.655		264. 154				29, 991	78. 50
MAMMA1000788	143.478		98. 983				29.032	46. 2
MAMMA 1000798	62, 822		139.860				11.942	32.5
MAMMA1000802	132.633	86. 328	341.638					61.50
								57.89
MAMMA1000810	150.779		372.241					10.7
MAMMA 10008 13	31.571	14.636					64, 626	62.0
AND DESIGNATION OF THE PARTY OF	197.602	134, 253	279.885					38.8
MAMMA1000814		A			1 1 4 4 1 1	1 29.208	31.671	
MAMMA1000824	65. 693		64.020				48 874	44 -
MAMMA1000824 MAMMA1000827	65. 693 146. 098	70.894	157. 448	47.656	39. 428	44. 524		
MAMMA1000824	65. 693	70.894 19.954	157. 448	47.656 13.557	39. 428 9. 407	44. 524 21. 580	16.602	6. 49 19. 27

Table 53

				-A-001 T	00 040 I	05 064 1	67 106	102 002
MAMMA1000839	157.898	138.792		13.084	86.648	85. 964	57. 386 13. 012	102. 963 20. 655
MANIMA 1000841	44. 843	37. 288		28. 351	19.319 48.610	37. 537	50, 804	35. 389
MAMMA1000842	174. 347	36.747		44. 926	2.185	78. 492 6. 547	5. 283	1.757
MAMMA 1000843	8. 643	4.650	14. 084	4. 758	15.747	23. 310	17. 728	15, 607
MASMA 1000845	40.044	33, 955		21.488		75. 853	98. 526	72.039
MAMMA 1000851	197.033	79. 321		96.446 17.429	73.025	33, 288	22. 320	21.685
MAMMA1000854	66.648	33. 221	63. 298	3.794	3. 995	2, 454	9, 158	3.568
MAMMA1000855	10. 264	4. 185	17.702 84.561	27.973	38.378	82.629	60. 529	25. 726
MAMMA1000856	186. 269	40. 945 121. 939	60.662	34. 958	42, 945	39. 557	20. 320	33. 032
WARMA1000859	64. 234	21.345	23, 693	16.808	28. 277	22.661	14. 238	14.693
MAMMA1000862 MAMMA1000863	40. 107 98. 576	70, 285	234. 996	67.796	55. 216	72. 466	36. 802	70.885
		0.000	0,000	0.000	2. 321	0.000	0,000	0.000
MAMMA 1000865 MAMMA 1000867	1. 106 46. 228	24.216	64. 376	21.736	17. 699	18. 758	10.742	6.964
MAMMA 1000875	124.814	80.537	231, 558	88.627	57.015	82. 859	46. 826	53.611
MAMMA1000876	87, 475	36. 523	94, 191	19.763	21. 465	42. 434	27. 201	24. 439
MAMMA 1000877	201.968	107, 716		164. 333	86.827	114. 380	80. 171	97.872
MAHMA1000878	99.671	67.833	257.022	71. 323	29.066	47. 487	36.714	37. 365
MAMMA1000878	76, 396	60.884	153, 335	45. 836	17.649	44. 996	19. 238	35, 353
MAMMA 1000881	63. 546	33.072	177. 731	43.034	30.410	31, 086	12. 184	38.045
MAMMA1000883	71.807	24, 931	43. 109	16.630	18.675	40. 320	44. 419	55. 440
MAMMA1000897	88. 466	0,000	7.404	0.000	0.000	0.000	0.000	0.721
MAMMA1000898	380.818	62.977	134.846	45. 311	63. 221	164. 332	122.071	52.933
MAMMA1000905	97. 555	63. 528	161, 117	57.777	42. 205	50.312	28. 216	42.710
MAMMA1000906	57.788	33.146	125.096	29,019	13.531	29. 380	16.982	14.930
MAMMA1000908	30.597	19.222	40.351	11.584	5. 445	10.392	13.469	11.612
MAMMA1000911	9. 952	29. 425	3.998	9.963	1.886	7.419	5. 350	126.400
MAMMA1000914	82.184	23. 137	69. 228	20.659	18.111	35, 329	22.616	18.859
MAMMA1000920	92.123	62.032	37. 206	16.675	15. 550	47. 235	47.680	26.80
MAMMA1000921	107.169	69.026		102, 347	60. 403	64.787	35. 902	77.42
MAMMA1000931	211.796	140. 234	424. 498	95.390	40. 229	51.643	49. 349	95. 21
MAMMA1000940	145. 411	82.982	268, 876	70. 972	55. 532	61. 420	51.119	60.32
MAMMA1000941	182.800	134.847		131.193	79.478	106.717	53. 292	91.18
MAMMA1000942	195.078	123, 131		117. 435	68. 234	90. 801	63. 506	75.81
MAMMA1000943	196. 926	99.988		109. 551	89.006	81.092 104.172	51.063 79.021	85.53 96.98
MAMMA1000952	161.019	97.081	355. 265	78, 330	98.779	41. 230	24. 471	5.89
MAMMA1000956	43.741	16.217	14. 918	11.103	5.840 42.610	47. 323	34. 337	45.56
MAMMA1000957	95. 532	53.066	225. 645	204. 962	120.611	123. 900	84. 354	140.99
MAMMA1000962 MAMMA1000966	281.600 151.087	192.048		111.282	64.746	81.685	51.694	78.95
MAMMA1000968	217. 975	107.043	313. 251	58. 469	41.964	45. 044	41.392	63.99
NAMMA1000972	18. 150	48.148	119. 482	22.427	18.041	15. 672	12.870	33.13
MAMMA1000973	36.667	18.879	24.787	11.758	12.527	19. 441	17.828	22.31
MAMMA1000975	44. 972	19.058	38. 995	20.137	30.793	22.854	65.817	45.39
MAMMA1000976	122.625	67.075	216. 981	70.671	50, 470	91.475	60.614	81.17
MAMMA1000979	81.812	102.452	145, 415	68. 435	53. 443	56. 902	38.749	89.75
MAMMA 1000986	118. 211	39. 368	239. 204	68.513	49, 208	56.431	42.354	94.15
MAMMA1000987	81.466	50.679	249.660	43.686	35. 580	49. 753	23.004	41.99
MAMMA1000988	150. 907	68, 191	242.562	63, 946	34. 252	81.162	48. 528	86.72
MAMMA 1000994	101. 984	21.000	41.248	21.154	26. 136	49. 152	44. 373	50.52
MAMMA1000998	166.669	75. 193	367, 111	91.202	105. 673	107.213	56.957	84.21
MAMMA1001003	73.580	37.252		47.279	34. 315	35. 674	26. 101	59.03
MAMMA1001007	3.055	0.000		0.000	1.411	3.633	0.800	0.00
MAMMA1001008	40. 892	31.048		38.501	74.831	38. 859	47. 979	31.17
	135. 486	126.855		93, 280	57. 270	56.674	44. 237 35. 346	52.3 16.6
MAMMA1001013	85. 681	25. 361		32.516	25. 227	20.809		29.7
MAMMA1001014		14 44 .		41.205	34. 542	34. 975	35. 352	36.7
MAMMA 1001014 MAMMA 1001021	93.867	49. 224			T A1 NEN	1 64 641	41 775	
MANMA1001014 MANMA1001021 MANMA1001024	93. 867 141. 736	49.918	229.735	52.670	41.069	54. 541	41.726	
MANNA 1001014 MANNA 1001021 MANNA 1001024 MANNA 1001025	93. 867 141. 736 13. 661	49.918 8.964	229. 735 12. 310	52.670 5.843	13.733	6.698	4. 305	5.0
MANMA1001014 MANMA1001021 MANMA1001024 MANMA1001025 MANMA1001028	93. 867 141. 736 13. 661 36. 353	49. 918 8. 964 24. 719	229. 735 12. 310 14. 061	52.670 5.843 10.363	13.733 34.518	6.698 16.233	4. 305 15. 746	5.0 11.3
MANNATOOTOT4 MANNATOOTO21 MANNATOOTO24 MANNATOOTO25 MANNATOOTO28 MANNATOOTO30	93.867 141.736 13.661 36.353 33.596	49. 918 8. 964 24. 719 27. 602	229.735 12.310 14.061 35.295	52.670 5.843 10.363 20.296	13.733 34.518 15.861	6. 698 16. 233 14. 989	4. 305 15. 746 25. 031	5. 0 11. 3 23. 5
MAMMATOOTOT4 MAMMATOOTO21 MAMMATOOTO24 MAMMATOOTO25 MAMMATOOTO28 MAMMATOOTO30 MAMMATOOTO30	93.867 141.736 13.661 36.353 33.596 235.880	49. 918 8. 964 24. 719 27. 602 125. 555	229.735 12.310 14.061 35.295 517.898	52.670 5.843 10.363 20.296 181.208	13.733 34.518 15.861 139.149	6. 698 16. 233 14. 989 129. 655	4. 305 15. 746 25. 031 96. 375	5. 09 11. 3 23. 53 134. 50
MANNA 1001021 MANNA 1001021 MANNA 1001024 MANNA 1001025 MANNA 1001028 MANNA 1001030	93.867 141.736 13.661 36.353 33.596	49. 918 8. 964 24. 719 27. 602	229.735 12.310 14.061 35.295 517.898 152.344	52.670 5.843 10.363 20.296	13.733 34.518 15.861 139.149	6. 698 16. 233 14. 989	4. 305 15. 746 25. 031	5. 09 11. 3 23. 53 134. 50 40. 93 51. 83

Table 54

AMANI 1001001 113,237 27, 502 43,846 32,708 45,924 47,820 46,929 16,614 AMANI 10010050 197,381 60,095 120,216 71,548 69,197 49,684 13,493 872 AMANI 1001054 102,456 62,728 134,003 63,234 43,343 11,184 33,007 39,478 AMANI 1001066 387,798 103,377 293,880 140,850 119,334 176,295 155,583 69,714 40,010 AMANI 1001066 387,798 103,377 293,880 140,850 119,334 176,295 155,583 69,714 40,010 AMANI 1001072 150,398 31,501 52,273 21,383 32,143 57,421 47,051 26,375 AMANI 1001072 150,398 31,501 52,273 21,383 32,143 57,421 47,051 26,375 AMANI 1001072 150,398 31,501 52,273 21,383 32,143 57,421 47,051 26,375 AMANI 1001074 104,201 41,827 240,332 94,124 56,071 89,717 16,387 14,966 AMANI 1001074 104,201 41,827 240,332 94,124 56,071 89,717 16,387 14,966 AMANI 1001075 32,081 34,601 23,705 29,752 21,195 21,195 21,184 14,757 17,497 AMANI 1001075 32,081 34,601 23,705 29,752 21,195 21,195 21,184 14,757 17,497 AMANI 1001075 30,781 31,780 21,78									
AMANICOTOTOS 218.483 23.847 68.163 22.306 10.499 41.046 45.779 31.087 AMANICOTOTOS 157.351 80.095 220.216 15.548 69.1997 48.684 13.493 49.872 AMANICOTOTOS 157.351 80.095 220.216 15.548 69.1997 48.684 13.493 49.872 AMANICOTOTOS 157.351 80.095 220.216 15.548 69.1997 48.684 13.493 49.872 AMANICOTOTOS 157.351 83.57 48.942 59.998 52.331 28.605 111.283 69.714 40.010 AMANICOTOTOS 31.357 48.942 79.998 52.331 28.605 111.283 69.714 40.010 AMANICOTOTOS 82.327 39.470 127.017 37.076 28.891 30.670 19.782 14.257 AMANICOTOTOS 150.398 31.601 52.273 21.983 32.143 57.421 47.051 52.5375 AMANICOTOTOS 150.398 31.601 52.273 21.983 32.143 57.421 47.051 52.5375 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.600 23.7728 00.099 17.502 43.163 36.606 61.41.739 265.507 AMANICOTOS 32.7728 00.000 37.728 00.000 17.502 43.163 36.606 61.41.739 265.507 AMANICOTOS 32.7728 00.000 37.728 00.000 17.502 43.163 36.606 61.41.739 265.507 AMANICOTOS 32.7728 00.000 37.728 00.000	IAMMA (001038	26. 248	12, 160	150.692	32.729	5. 309	10.436	15. 903	27. 263
AMANICOTOTOS 218.483 23.847 68.163 22.306 10.499 41.046 45.779 31.087 AMANICOTOTOS 157.351 80.095 220.216 15.548 69.1997 48.684 13.493 49.872 AMANICOTOTOS 157.351 80.095 220.216 15.548 69.1997 48.684 13.493 49.872 AMANICOTOTOS 157.351 80.095 220.216 15.548 69.1997 48.684 13.493 49.872 AMANICOTOTOS 157.351 83.57 48.942 59.998 52.331 28.605 111.283 69.714 40.010 AMANICOTOTOS 31.357 48.942 79.998 52.331 28.605 111.283 69.714 40.010 AMANICOTOTOS 82.327 39.470 127.017 37.076 28.891 30.670 19.782 14.257 AMANICOTOTOS 150.398 31.601 52.273 21.983 32.143 57.421 47.051 52.5375 AMANICOTOTOS 150.398 31.601 52.273 21.983 32.143 57.421 47.051 52.5375 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.601 23.705 29.782 21.196 23.184 14.757 17.497 AMANICOTOTOS 32.081 36.600 23.7728 00.099 17.502 43.163 36.606 61.41.739 265.507 AMANICOTOS 32.7728 00.000 37.728 00.000 17.502 43.163 36.606 61.41.739 265.507 AMANICOTOS 32.7728 00.000 37.728 00.000 17.502 43.163 36.606 61.41.739 265.507 AMANICOTOS 32.7728 00.000 37.728 00.000	MAMMA 1001041	113 237	27, 602	43.846	32,708	45. 924	47.820	46, 929	16.614
AMANA 1001050 157.351 60.095 220.216 71.548 69.197 49.684 13.493 49.872 AMANA 1001058 136.357 48.942 59.998 52.931 26.661 111.283 69.714 40.010 40.01068 387.798 10.377 283.890 140.850 119.334 111.283 69.714 40.010 40.01066 387.798 10.377 283.890 140.850 119.334 111.283 69.714 40.010 40.01067 20.377 293.890 140.850 119.334 117.295 156.853 60.324 40.010 40.01067 20.377 293.890 140.850 119.334 17.625 156.853 60.324 40.010 40.01067 20.377 293.890 140.850 119.334 17.625 156.853 60.324 47.051 26.375 40.001072 150.398 31.601 52.273 17.803 32.181 37.217 11.406 43.228 24.053 24.142 55.775 40.0010074 104.201 41.827 240.332 94.124 55.071 89.717 15.387 14.966 40.001078 102.185 111.402 317.478 75.869 35.841 49.660 67.285 67.244 40.01001078 102.185 111.402 317.478 75.869 35.841 49.660 67.285 67.244 40.01001078 102.185 111.402 317.478 75.869 35.841 49.660 67.285 67.244 40.01001078 102.185 111.402 317.478 75.869 35.841 49.660 67.285 67.244 40.01001078 102.185 111.402 27.525 95.003 81.992 186.406 141.793 765.507 40.001001078 102.185 111.402 27.525 95.003 81.992 186.406 141.793 765.507 40.001001078 102.185 111.402 27.525 05.000 18.321 4.553 00.000 0.000 0.0000 40.0000 40.0000 38.7248 21.740 27.506 112.379 42.165 15.31 8.849 11.524 4.155 40.001001078 33.870 77.506 112.379 42.165 75.385 80.000 40.000 0.000 0.0000 40.00000 40.0000 40.0000 40.0000 40.0000 40.0000 40									
ALMAN 1001068 102, 455 62, 728 134, 003 63, 324 43, 343 221, 184 38, 007 39, 478 ALMAN 1001068 387, 788 103, 377 293, 899 87, 823 12, 60, 61 117, 283 69, 714 40, 001 ALMAN 1001068 387, 788 103, 377 293, 890 40, 850 119, 314 176, 295 158, 553 60, 324 ALMAN 1001067 82, 327 39, 420 127, 517 31, 516 32, 218 17, 217 11, 406 43, 228 24, 655 124, 47, 651 765, 375 ALMAN 1001073 101, 957 23, 218 17, 217 11, 406 43, 228 24, 655 124, 42, 51, 766 ALMAN 1001073 101, 957 23, 218 17, 217 11, 406 43, 228 24, 655 124, 147, 517 124, 318 174, 314 174, 327 14, 327									
ALBEAL MODITORS 155, 357 48, 947 59, 998 52, 931 26, 061 117, 283 69, 714 40, 010 ALBEAL MODITORS 38, 387 78, 103, 377 293, 859 140, 850 118, 354 116, 295 158, 585 06, 324 ALBEAL MODITORS 38, 387 39, 420 127, 017 37, 076 29, 891 30, 670 19, 782 14, 257 ALBEAL MODITORS 398 31, 6501 52, 273 21, 983 32, 143 57, 421 47, 051 26, 375 ALBEAL MODITORS 398 31, 6501 52, 273 21, 983 22, 143 57, 421 47, 051 26, 375 ALBEAL MODITORS 30, 398 31, 6501 52, 273 21, 983 22, 143 57, 421 47, 051 26, 375 ALBEAL MODITORS 30, 398 31, 6501 52, 273 21, 983 22, 143 57, 421 47, 051 26, 375 ALBEAL MODITORS 30, 983 31, 6501 52, 273 21, 983 22, 143 57, 421 47, 051 26, 375 ALBEAL MODITORS 32, 081 34, 6501 23, 376 29, 378 22, 11, 96 43, 228 24, 053 24, 142 57, 144 45, 577 ALBEAL MODITORS 32, 081 31, 6501 23, 376 29, 383 38, 982 21, 196 23, 184 41, 757 17, 96 ALBEAL MODITORS 30, 2185 111, 402 317, 478 75, 869 35, 841 49, 650 67, 285 67, 244 ALBEAL MODITORS 37, 248 21, 754 11, 022 317, 478 75, 869 35, 841 49, 650 67, 285 67, 244 ALBEAL MODITORS 30, 254 39, 773 20, 039 77, 502 43, 163 25, 358 17, 452 14, 352 ALBEAL MODITORS 30, 254 25, 308 48, 577 16, 425 15, 153 18, 849 11, 524 41, 55 ALBEAL MODITORS 30, 554 25, 308 48, 577 16, 425 15, 153 18, 849 11, 524 41, 55 ALBEAL MODITORS 30, 554 25, 308 48, 577 16, 425 15, 153 18, 849 11, 524 41, 155 ALBEAL MODITORS 30, 574 223, 060 633, 480 194, 265 80, 590 45, 156 18, 891 45, 540 46,									
AMMALIODIOSE 387,798 103,377 293,890 100,850 119,334 176,295 158,553 60,324 AMMALIODIOST 82,327 39,420 127,017 37,076 29,891 30,670 19,762 14,257 AMMALIODIOTZ 150,388 31.501 52,273 21,983 32,143 57,421 47,051 25,375 AMMALIODIOTZ 150,388 31.501 52,273 21,983 32,143 57,421 47,051 25,375 AMMALIODIOTZ 104,201 41,827 240,332 94,124 55,071 89,717 16,387 14,167 AMMALIODIOTS 12,081 34,601 23,705 29,782 21,196 23,184 14,757 17,497 AMMALIODIOTS 102,185 111,402 317,478 75,889 35,841 49,560 67,285 57,285 11,402 AMMALIODIOTS 102,185 111,402 317,478 75,889 35,841 49,560 67,285 57,285 24,444 4,757 17,497 AMMALIODIOTS 27,262 210,764 130,259 85,003 81,982 185,606 141,739 265,507 AMMALIODIOTS 37,578 11,402 37,572 0,009 17,502 43,163 25,358 17,452 14,352 AMMALIODIOTS 37,578 11,403 27,522 0,000 18,321 4,593 0,000 0,000 AMMALIODIOTS 37,578 11,403 27,572 0,000 18,321 4,593 0,000 0,000 AMMALIODIOTS 37,578 11,403 27,572 0,000 18,321 4,593 0,000 0,000 AMMALIODIOTS 37,578 11,403 27,572 0,000 18,321 4,593 0,000 0,000 AMMALIODIOTS 38,777 111,266 113,121 52,426 80,960 45,158 18,849 11,524 4,155 AMMALIODIOTS 18,777 111,266 113,121 52,426 80,960 45,158 16,891 45,552 AMMALIODIOTS 18,777 11,267 113,121 52,426 80,960 45,158 16,891 45,552 AMMALIODIOTS 18,777 11,784 77,89 60 443,796 9,930 5,930 5,930 45,158 16,891 45,552 AMMALIODIOTS 18,777 11,784 7,785 88 67,249 59,919 67,023 43,805 3,168 43,804 11,574 47,896 47,916	KAMNA 1001054								
AMMARIOTORY 32, 377 39, 470 127, 293, 390 140, 350 119, 334 176, 295 158, 553 60, 324 AMMARIOTORY 32, 327 39, 420 127, 017 37, 076 29, 391 30, 670 19, 782 14, 254 AMMARIOTORY 150, 338 31, 501 52, 273 21, 983 32, 143 57, 421 47, 051 26, 375 AMBARIOTORY 104, 201 41, 827 240, 332 94, 124 55, 071 39, 717 16, 387 14, 584 AMBARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMBARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMBARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMBARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMBARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMMARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMMARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMMARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 23, 184 14, 757 17, 497 AMMARIOTORY 32, 081 34, 601 23, 705 29, 782 71, 196 24, 163 26, 358 17, 452 14, 197 AMMARIOTORY 34, 35, 787 11, 403 27, 522 0, 000 18, 321 4, 593 0, 000 0, 0, 000 AMBARIOTORY 35, 78, 78, 78, 78, 78, 78, 78, 78, 78, 78	AMMA 1001059	136.357	48. 942	59.998	52.931	26.061	111. 283	69.714	40.010
AMMALTODIOST 82, 327 39, 420 127, 017 37, 076 29,891 30,670 19,782 41,257 AMMALTODIOST 150,389 31,501 52,273 11,806 43,228 24,053 24,142 5,176 AMMALTODIOST 150,389 31,501 52,273 11,806 43,228 24,053 24,142 5,176 AMMALTODIOST 101,957 23,218 17,217 11,406 43,228 24,053 24,142 5,176 AMMALTODIOST 32,081 34,601 23,705 29,782 21,195 23,184 14,737 17,497 AMMALTODIOST 32,081 34,601 23,705 29,782 21,195 23,184 14,737 17,497 AMMALTODIOST 30,285 111,402 317,478 75,869 35,841 49,560 67,285 67,244 AMMALTODIOST 30,728 120,784 130,259 83,003 81,892 186,406 11,739 265,567 AMMALTODIOST 30,284 39,773 20,039 17,502 43,163 26,358 17,452 14,152 AMMALTODIOST 30,554 29,306 48,577 16,425 15,153 18,469 11,524 4,155 AMMALTODIOST 30,554 25,306 48,577 16,425 17,562 43,163 26,358 17,452 14,152 AMMALTODIOST 30,554 25,306 48,577 16,425 17,562 43,163 30,358 11,824 62,344 AMMALTODIOST 35,186 77,565 11,123 11,123 82,426 80,960 45,158 16,891 11,524 4,155 AMMALTODIOST 35,187 11,1264 113,121 82,426 80,960 45,158 16,891 45,652 AMMALTODIOST 30,273 80,003 81,982 11,524 4,155 AMMALTODIOST 30,273 80,003 81,982 11,524 4,155 AMMALTODIOST 30,503 81,982 81,	AAMMA1001066	387, 798	103, 377	293, 890	140.850	119.334	176.295	158. 563	60.324
MANIALOTOTOT 150, 338 31, 501 52, 273 21, 938 32, 143 57, 421 47, 051 26, 375						29 891		19.782	
ALBERT 1001074									
ALBEAN A									
LANMAR 1001075 32.081 34.601 23.705 29.782 21.196 23.184 14.757 17.497 17.497 17.497 17.497 102.185 111.402 117.478 75.889 35.841 49.660 67.285 67.244 14.1861 11.1861									
NAMBA 10011078 102: 185 111. 402 117. 478 75. 869 15. 841 49. 660 67. 285 67. 244 AANAHA 1001080 367. 248 210.764 130. 259 83. 033 81. 982 186. 406 141. 739 265. 507 AANBA 1001082 50. 524 39. 773 20. 039 17. 602 43. 163 26. 338 17. 452 14. 1352 AANBA 1001081 3. 576 11. 403 27. 522 0. 000 18. 321 4. 593 0. 000 0. 0. 000 AANBA 1001092 50. 554 25. 306 48. 577 16. 4. 425 15. 153 18. 849 11. 524 4. 155 AANBA 1001094 53. 180 72. 506 112. 379 42. 145 18. 348 19. 1524 4. 155 AANBA 1001095 138. 777 111. 2261 113. 121 82. 425 80. 960 45. 158 16. 831 45. 652 AANBA 1001105 138. 777 111. 2261 113. 121 82. 425 80. 960 45. 158 16. 831 45. 652 AANBA 1001110 135. 141 8. 661 7. 407 3. 823 5. 537 6. 280 3. 216 4. 392 AANBA 1001121 32 99. 120 223. 660 683. 480 194. 522 164. 920 119. 375 96. 413 88. 784 AANBA 1001123 293. 126 87. 724 529. 603 144. 407 119. 311 11. 573 67. 515 94. 605 AANBA 1001139 291. 212 867. 784 447. 960 473. 187 227. 579 348. 627 121. 382 173. 640 AANBA 1001139 291. 212 867. 784 447. 960 473. 187 227. 579 348. 627 121. 382 173. 640 AANBA 1001143 163. 308 70. 387 153. 588 67. 249 59. 919 67. 023 43. 805 40. 903 AANBA 1001150 80. 076 29. 005 50. 289 15. 249 7. 495 33. 674 48. 052 22. 62. 277 16. 337 13. 966 AANBA 1001151 38. 076 29. 005 50. 289 15. 249 7. 495 33. 674 48. 052 22. 62. 278 AANBA 1001159 40. 247 28. 783 19. 301 13. 704 8. 444 23. 464 24. 48. 48. 48. 48. 48. 48. 48. 48. 48. 4	MANNA1001074	104. 201	41.827						
MANIMATIODI 168	MAMMA1001075	32.081	34, 601	23.705	29.782	21, 196			17. 497
MANMATODITOR 367, 248 210, 764 130, 259 83, 003 81, 982 186, 406 141, 739 256, 507	MAMMA1001078	102, 185	111, 402	317, 478	75, 869	35, 841	49.660	67. 285	67. 244
MAINAI (1001082 SO, 264 39.773 20.039 17.602 43.163 26.358 17.452 14.152 MAINAI (1001091 3.575 11.403 27.522 0.000 18.321 4.593 0.000 0.000 MANNAI (1001092 50.554 25.306 48.577 16.425 15.153 18.849 11.524 4.155 MAINAI (1001094 333, 180 72.506 112.379 42.145 78.386 130.368 113.824 62.564 MAINAI (1001095 138.777 111.2261 113.121 82.426 80.960 45.158 16.891 45.652 MAINAI (1001096 138.777 111.2261 113.121 82.426 80.960 45.158 16.891 45.652 MAINAI (1001106 15.141 8.661 7.407 3.823 5.537 6.280 3.216 4.392 MAINAI (1001126 299.120 223.066 883.480 194.592 164.920 193.75 96.418 8.784 MAINAI (1001139 279.1212 867.784 447.960 473.187 227.579 348.627 121.382 173.640 MAINAI (1001139 231.212 867.784 447.960 473.187 227.579 348.627 121.382 173.640 MAINAI (1001141 36.320 18.295 40.066 9.930 5.020 26.277 16.331 13.966 MAINAI (1001139 13.343.866 187.024 529.603 144.907 119.301 111.573 67.515 94.605 MAINAI (1001141 36.3308 70.387 153.588 67.249 59.919 67.023 43.805 40.903 MAINAI (1001141 36.308 70.387 153.588 67.249 59.919 67.023 43.805 40.903 MAINAI (1001145 110.718 43.148 141.067 0.850 11.851 11.000 10.119 13.322 MAINAI (1001159 80.076 29.005 50.289 15.249 7.495 33.674 48.052 22.259 MAINAI (1001159 46.847 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MAINAI (1001159 46.847 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.404 21.664 24.248 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.004 21.664 24.248 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.004 21.664 24.248 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.004 21.664 24.248 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.007 107.154 96.161 79.043 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.007 107.154 96.161 79.043 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.007 107.154 96.161 79.043 MAINAI (1001159 46.847 28.763 19.301 13.704 8.446 23.304 21.664 24.248 MAINAI (1001159 46.847 28.765 19.301 13.704 8.466 27.301 13.600 13.600 13.600 13.600 13.600 13.600 13.			210 764						
MAIMAI 10011991 3.576									
MANIMA 10011092									
MAINMAI DOITIOG 138.777 111.2281 113.121 82.425 80.960 45.158 11.824 62.964 MAINMAI DOITIOS 138.777 111.2281 113.121 82.425 80.960 45.158 16.837 45.652 MAINMAI DOITIOS 15.141 8.661 7.407 3.823 5.537 6.280 3.216 4.392 MAINMAI DOITIOS 199.120 223.060 683.480 194.522 164.920 119.375 96.413 88.784 MAINMAI DOITIGS 299.120 23.060 683.480 194.522 164.920 119.375 96.413 88.784 140.001133 243.826 187.024 829.603 144.907 119.301 111.573 67.515 94.605 MAINMAI DOITIGS 29.121 887.784 447.960 473.187 277.579 348.627 121.382 173.640 MAINMAI DOITIGS 183.20 18.295 40.066 9.930 5.202 265.277 16.337 13.996 MAINMAI DOITIGS 163.308 70.387 153.588 67.249 59.919 57.023 43.805 40.903 MAINMAI DOITIGS 80.076 29.005 50.289 18.249 7.495 33.674 48.052 22.629 MAINMAI DOITIGS 80.076 29.005 50.289 18.249 7.495 33.674 48.052 22.629 MAINMAI DOITIGS 80.076 29.005 50.289 18.249 7.495 33.674 48.052 22.629 MAINMAI DOITIGS 80.076 29.005 50.289 18.249 7.495 33.674 48.052 22.629 MAINMAI DOITIGS 80.076 29.005 50.289 18.249 7.495 33.674 48.052 22.629 MAINMAI DOITIGS 80.076 123.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.601 233.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.601 233.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.601 233.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.601 233.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.299 51.198 67.587 29.962 40.684 78.949 43.247 18.714 MAINMAI DOITIGS 185.601 233.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.201 33.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.201 33.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.201 33.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.201 33.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.201 33.229 485.605 141.151 109.607 107.154 95.161 79.043 MAINMAI DOITIGS 185.518 85.200 33.506 69.537 59.765 53.99 50.065 53.806 141.151 107.158 141.159 1									
MAINIA 1001105	MAMMA 1001092				16.425				
MAINMATODITIOS 138.777 111.2261 113.121 82.426 80.960 45.158 16.891 45.552 MAINMATODITIO 15.141 8.661 7.407 3.823 5.537 6.280 3.216 4.992 118.001110 15.141 8.661 7.407 3.823 5.537 6.280 3.216 4.992 118.001110 15.141 8.661 7.407 3.823 5.537 6.280 3.216 4.992 119.375 95.413 88.784 MAINMATODITION 243.826 187.024 529.603 144.907 119.301 111.573 67.515 94.605 MAINMATODITION 243.826 187.024 529.603 144.907 119.301 111.573 67.515 94.605 MAINMATODITION 25.800 18.295 40.066 9.930 5.202 26.277 16.337 13.996 MAINMATODITION 153.308 70.387 153.588 67.244 59.919 67.023 43.805 40.903 MAINMATODITION 153.308 70.387 153.588 67.244 59.919 67.023 43.805 40.903 MAINMATODITION 154 43.148 141.067 30.890 31.851 11.000 10.119 13.122 MAINMATODITION 155 80.076 29.005 50.289 15.249 7.495 33.674 48.052 22.529 MAINMATODITION 155 80.076 29.005 50.289 15.249 7.495 33.674 48.052 22.529 MAINMATODITION 155 48.647 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MAINMATODITION 159 45.847 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MAINMATODITION 159 45.847 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MAINMATODITION 159 45.857 35.688 88.127 33.728 40.701 41.280 16.749 26.312 MAINMATODITION 159 55.118 85.120 303.506 69.532 51.017 85.296 42.211 48.082 MAINMATODITION 159 56.567 31.052 16.618 30.386 22.337 29.809 50.065 54.044 MAINMATODITION 198 60.567 31.052 16.618 30.386 22.337 29.809 50.065 54.044 MAINMATODITION 198 22.9338 561.555 755.244 695.028 15.017 85.296 42.211 48.082 MAINMATODITION 198 22.9338 561.555 755.244 695.028 21.661 39.122 50.157 24.854 664.569 248.672 218.550 168.136 144.829 179.567 MAINMATODITION 198 22.9338 561.555 755.244 695.028 20.861 53.6623 412.766 746.035 MAINMATODITION 198 22.9338 561.555 755.244 695.028 695.028 60.874 42.217 20.800 170.551 10.121 303.059 85.243 72.915 53.330 44.564 52.183 MAINMATODITION 198 22.9338 561.555 755.245 695.028 60.874 42.345 60.874 42.345 60.874 42.345 60.874 42.345 60.874 42.345 60.874 64.935 77.955 67.054 64.855 77.955 77.055 77.055 77.055 77.055 77.055 77.055 77.055 77.055 77.055 77.05	MANNA1001094	353, 180	72.506	112.379	42.145	78.386	130.358	113.824	62.964
MAMMATODI 110 15. 141 8. 661 7. 407 3. 823 5.537 6. 280 3. 216 4. 392 MAMMATODI 126 299. 120 223. 060 683.480 194. 522 164. 920 119. 375 95. 413 88. 784 MAMMATODI 133 243. 826 187. 024 529. 603 144. 907 119. 301 111. 573 67. 515 94. 605 MAMMATODI 139 291. 212 267. 784 447. 960 473. 187 227. 579 348. 627 121. 382 173. 640 MAMMATODI 141 35. 320 18. 295 40. 066 9. 930 5. 202 26. 277 16. 337 13. 966 MAMMATODI 143 163. 308 70. 387 153. 588 67. 249 59. 919 67. 023 43. 805 40. 903 MAMMATODI 150 80. 076 29. 005 50. 289 15. 249 59. 919 67. 023 43. 805 40. 903 MAMMATODI 150 80. 076 29. 005 50. 289 15. 249 7. 495 33. 674 48. 052 22. 529 MAMMATODI 150 80. 076 29. 005 50. 289 15. 249 7. 495 33. 674 48. 052 22. 529 MAMMATODI 151 85. 601 233. 229 435. 605 141. 151 109. 607 107. 154 95. 161 79. 043 MAMMATODI 161 185. 601 233. 229 435. 605 141. 151 109. 607 107. 154 95. 161 79. 043 MAMMATODI 162 196. 799 51. 198 67. 587 29. 962 40. 884 78. 949 43. 247 48. 742 MAMMATODI 188 155. 118 85. 120 303. 506 89. 532 51. 017 85. 296 42. 211 48. 082 MAMMATODI 188 155. 118 85. 120 303. 506 89. 532 51. 017 85. 296 42. 211 48. 082 MAMMATODI 189 60. 587 31. 052 16. 618 30. 386 22. 337 29. 809 50. 065 54. 044 MAMMATODI 189 60. 587 31. 052 16. 618 30. 386 22. 337 29. 809 50. 065 54. 044 MAMMATODI 189 60. 587 31. 052 16. 618 30. 386 22. 337 29. 809 50. 065 54. 044 MAMMATODI 189 60. 587 31. 052 16. 618 30. 386 22. 337 29. 809 50. 065 54. 044 MAMMATODI 189 60. 587 31. 052 16. 618 30. 386 22. 337 29. 809 50. 065 54. 044 MAMMATODI 189 60. 587 31. 052 16. 618 30. 386 22. 337 29. 809 50. 065 54. 044 MAMMATODI 205 33. 103 114. 504 202. 256 55. 195 77. 217 61. 327 43. 601 43. 601 MAM							45, 158	16, 891	45, 652
MARIA 1001126 299.120 223.060 683.480 194.522 164.920 119.375 95.413 88.784 MARIA 1001139 291.212 867.784 447.960 473.187 227.579 348.627 121.382 173.640 MARIA 1001139 291.212 867.784 447.960 473.187 227.579 348.627 121.382 173.640 MARIA 1001141 35.320 18.295 40.066 9.930 5.202 26.277 16.337 13.996 MARIA 1001143 163.308 70.387 153.588 67.249 59.919 67.023 43.805 40.903 MARIA 1001143 110.718 43.148 141.067 30.890 31.851 11.000 10.119 33.122 MARIA 1001150 80.076 29.005 50.289 15.249 7.495 33.674 48.052 22.529 MARIA 1001153 46.447 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MARIA 1001159 46.847 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MARIA 1001161 185.501 233.229 485.605 141.151 109.507 107.154 95.161 79.043 MARIA 1001161 185.505 33.688 88.127 33.728 40.701 41.280 15.749 26.312 MARIA 1001181 116.505 35.688 88.127 33.728 40.701 41.280 15.749 26.312 MARIA 1001181 10.505 35.688 88.127 33.728 40.701 41.280 15.749 26.312 MARIA 1001181 10.505 35.688 88.127 33.728 40.701 41.280 15.749 26.312 MARIA 1001183 15.5118 85.120 303.506 69.532 51.017 85.296 42.211 48.082									
MAMMATODIT33									
MAMMA1001139 291.212 867.784 447.960 473.187 227.579 348.627 121.382 173.640									
MAIMATIOD 141 36.320 18.295 40.066 9.930 5.202 26.277 16.337 13.996									
MAMMATIOD	MAMMA1001139		867.784			227. 579			
MAMMA 1001 143 163 308 70 387 153 588 67 249 59 91 97 023 43 805 40 903 MAMMA 1001 143 163 308 70 387 153 588 67 249 70 30 890 31 851 11 000 10 11 31 32 22 MAMMA 1001 150 80 076 29 005 50 289 15 249 7 495 33 674 48 052 22 52 59 MAMMA 1001 154 203 206 129 777 429 878 121 700 90 014 77 333 45 155 71 154 MAMMA 1001 154 203 206 123 777 429 878 121 700 90 014 77 333 45 155 71 154 MAMMA 1001 156 85 601 233 229 485 605 141 151 109 607 107 154 96 161 79 043 MAMMA 1001 162 196 299 51 198 87 587 29 962 40 884 78 894 43 247 18 714 MAMMA 1001 18 155 18 85 120 303 506 69 532 51 017 55 295 42 21 48 082 MAMMA 1001 18 155 18 85 120 303 506 69 532 51 017 55 295 42 21 48 082 MAMMA 1001 189 50 587 31 052 16 618 30 386 22 337 29 809 50 065 54 044 MAMMA 1001 189 25 18 033 41 909 22 249 21 661 39 122 50 157 24 623 MAMMA 1001 190 120 521 18 033 41 909 22 249 21 661 39 122 50 157 24 623 MAMMA 1001 190 120 521 18 033 590 85 243 72 915 53 390 44 564 52 183 MAMMA 1001 190 120 120 131 14 504 202 256 55 195 71 217 61 327 43 601 48 988 MAMMA 1001 203 170 551 101 121 330 590 85 243 72 915 53 390 44 564 52 183 MAMMA 1001 120 132 103 114 504 202 256 55 95 77 217 61 327 43 601 48 988 MAMMA 1001 120 123 133 154 557 404 346 110 968 91 387 74 974 36 60 48 988 MAMMA 1001 123 123 103 14 504 202 256 55 95 77 217 218 20 316 32 44 48 99	MAMMA1001141	36.320	18. 295	40.066	9. 930	5. 202	26.277	16.337	13.996
MAIMATIOD1145	MAMMA1001143	163.308		153, 588	67, 249	59, 919	67.023	43.805	40, 903
MAIMAT1001150 80.076 29.005 50.289 15.249 7.495 33.674 48.052 22.629 MAIMAT1001154 203.206 129.777 429.878 121.700 90.014 77.333 45.155 71.154 MAIMAT1001159 46.847 28.763 19.301 13.704 8.444 23.404 21.664 24.248 MAIMAT1001161 185.601 233.229 485.605 141.151 109.607 107.154 96.161 79.043 MAIMAT1001162 196.299 51.198 67.587 29.962 40.684 78.949 43.247 18.714 MAIMAT1001181 116.505 35.688 88.127 33.728 40.701 41.280 42.248 43.404 43.									
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MANIMATIODITS9									
MAIMMATODITES 185.601 233.229 485.605 141.151 109.607 107.154 96.161 79.043									
MARMATOOT162	MAMMA1001159								
MANIMATODIT81 196.299 51.198 67.587 29.962 40.684 78.949 43.247 18.714	MANMA 1001161	185.601	233.229	485. 605	141. 151		107.154	96. 161	79.043
MAIMA 1001 18	MANNA 1001162		51, 198	67, 587	29, 962	40, 584	78, 949	43, 247	18,714
NAMMA1001188								16.749	26.312
MANNAIOO1189 60.587 31.052 16.618 30.386 22.337 29.809 50.065 54.044									
MANNA 1001191 120.521 18.093 41.909 22.249 21.661 39.122 50.157 24.623 MANNA 1001198 229.338 561.556 755.924 695.028 205.811 536.623 412.766 746.035 746.0									
MANNA 1001198 229.338 561.556 755.924 695.028 205.811 536.623 412.766 746.035 MANNA 1001203 322.950 274.854 664.559 248.672 218.550 168.136 144.829 179.567 MANNA 1001203 170.551 101.121 330.599 85.243 72.915 53.390 44.564 52.183 MANNA 1001206 32.103 114.504 202.256 65.195 71.217 61.327 43.601 48.988 MANNA 1001208 55.417 28.101 30.608 21.282 25.686 27.394 20.015 15.433 MANNA 1001215 199.721 123.016 194.852 82.919 72.839 87.841 68.245 60.078 MANNA 1001220 223.133 154.557 404.346 110.968 91.387 74.073 58.534 62.841 MANNA 1001222 5.585 4.936 6.763 1.952 0.474 2.171 20.800 5.022 MANNA 1001223 94.809 29.294 42.345 15.601 20.861 20.316 32.446 15.726 MANNA 1001232 130.199 45.692 227.125 47.671 38.837 45.692 59.906 32.862 MANNA 1001237 29.560 11.083 23.224 7.241 4.489 20.199 16.883 11.003 MANNA 1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 MANNA 1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA 1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA 1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA 1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA 1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA 1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.566 MANNA 1001250 154.426 54.153 81.155 52.438 46.566 80.874 64.937 87.761 MANNA 1001260 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANNA 1001260 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANNA 1001260 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANNA 1									
MANIMATOD1202 322.950 274.854 664.569 248.672 218.550 168.136 144.829 179.567									
MANNATODIZOS 170.551 101.121 330.599 85.243 72.915 53.390 44.564 52.183 MANNATODIZOS 132.103 114.504 202.256 65.195 71.217 61.327 43.601 48.988 MANNATODIZOS 55.417 28.101 30.608 21.282 25.686 27.394 20.016 15.433 MANNATODIZOS 223.133 154.557 404.346 110.968 91.387 72.839 87.841 68.245 60.078 MANNATODIZZO 223.133 154.557 404.346 110.968 91.387 74.073 58.534 62.841 MANNATODIZZO 5.585 4.936 6.763 1.952 0.474 2.171 20.800 5.022 MANNATODIZZO 394.809 29.294 42.345 15.601 20.861 20.316 32.446 15.726 MANNATODIZZO 130.199 45.692 227.125 47.671 38.837 45.692 59.906 32.862 MANNATODIZZO 130.199 45.692 227.125 47.671 38.837 45.692 59.906 32.862 MANNATODIZZO 20.832 130.199 45.692 227.125 47.671 38.837 45.692 59.906 32.862 MANNATODIZZO 20.832 11.598 47.127 7.253 32.688 20.073 7.954 6.544 MANNATODIZZO 20.832 11.598 47.127 7.253 32.688 20.073 7.954 6.544 MANNATODIZZO 20.832 11.598 47.127 7.253 32.686 20.073 7.954 6.544 MANNATODIZZO 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNATODIZZO 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNATODIZZO 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNATODIZZO 53.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNATODIZZO 53.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNATODIZZO 53.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNATODIZZO 53.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNATODIZZO 53.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNATODIZZO 53.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNATODIZZO 53.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 MANNATODIZZO 53.434 60.199 204.903 48.739 63.272 100.485 93.658 72.201 MANNATODIZZO 153.434 60.199 204.903 48.739 63.272 100.485 93.658 73.658 34.076 53.899 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANNATODIZZO 1528 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MANNATODIZZO 168 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MANNATODIZZO 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 MANNATODIZZO	MAMMA1001198	229.338	561.556	755. 924					
MANNA1001205 132.103 114.504 202.256 55.195 71.217 61.327 43.601 48.988 MANNA1001208 55.417 28.101 30.608 21.282 25.686 27.394 20.016 15.433 MANNA1001215 199.721 123.016 194.852 82.919 72.839 87.841 68.245 60.078 MANNA1001220 223.133 154.557 404.346 110.968 91.387 74.073 58.534 62.841 MANNA1001222 5.585 4.936 6.763 1.952 0.474 2.171 20.800 5.022 MANNA1001223 94.809 29.294 42.345 15.601 20.861 20.316 32.446 15.726 MANNA1001232 130.199 45.692 227.125 47.671 38.837 45.692 59.906 32.862 MANNA1001234 129.344 27.935 227.692 95.815 64.344 61.799 49.210 34.673 MANNA1001234 129.344 27.935 227.692 95.815 64.344 61.799 49.210 34.673 MANNA1001234 20.832 11.598 47.127 7.253 32.689 20.073 7.954 6.544 MANNA1001243 20.832 11.598 47.127 7.253 32.689 20.073 7.954 6.544 MANNA1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 MANNA1001244 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 25.335 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.355 25.	MAMMA1001202	322.950	274.854	664.569	248.672	218.550	168.136	144.829	179.567
MANNA1001205 132.103 114.504 202.256 65.195 71.217 61.327 43.601 48.988 MANNA1001208 55.417 28.101 30.608 21.282 25.686 27.394 20.016 15.433 MANNA1001215 199.721 123.016 194.852 82.919 72.839 87.841 68.245 60.078 MANNA1001220 223.133 154.557 404.346 110.968 91.387 74.073 58.534 62.841 MANNA1001222 5.585 4.936 6.763 1.952 0.474 2.171 20.800 5.022 MANNA1001223 94.809 29.294 42.345 15.601 20.861 20.316 32.446 15.726 MANNA1001232 130.199 45.692 227.125 47.671 38.837 45.692 59.906 32.862 MANNA1001234 129.344 27.935 227.692 95.815 64.344 61.799 49.210 34.673 MANNA1001234 129.344 27.935 227.692 95.815 64.344 61.799 49.210 34.673 MANNA1001234 20.832 11.083 23.224 7.241 4.489 20.199 16.883 11.003 MANNA1001243 20.832 11.598 47.127 7.253 32.689 20.073 7.954 6.544 MANNA1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 MANNA1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA1001256 169.303 81.917 266.686 187.649 131.656 44.850 55.325 59.866 MANNA1001250 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNA1001260 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 MANNA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNA1001260 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 MANNA1001260 55.325 59.866 MANN	MANUATO01203			330.599	85, 243	72, 915	53, 390	44. 564	52, 183
MANNA1001215									
NABIDIA 1001215 199.721 123.016 194.852 82.919 72.839 87.841 68.245 60.078 NABIDIA 1001220 223.133 154.557 404.346 110.968 91.387 74.073 58.534 62.841 NABIDIA 1001222 5.585 4.936 6.763 1.952 0.474 2.171 20.800 5.022 NABIDIA 1001223 94.809 29.294 42.345 15.601 20.861 20.316 32.446 15.726 NABIDIA 1001234 129.3344 27.935 227.125 47.671 38.837 45.692 59.906 32.862 NABIDIA 1001237 29.560 11.083 23.224 7.241 4.489 20.199 16.883 11.003 NABIDIA 1001237 29.560 11.083 23.224 7.241 4.489 20.199 16.883 11.003 NABIDIA 1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 NABIDIA 1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 NABIDIA 1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 NABIDIA 1001256 169.303 81.917 266.686 87.649 131.656 44.850 55.325 59.786 NABIDIA 1001250 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 NABIDIA 1001260 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 NABIDIA 1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 NABIDIA 1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 NABIDIA 1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 NABIDIA 1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.668 76.590 NABIDIA 1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.668 76.590 NABIDIA 1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.668 76.590 NABIDIA 1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.668 76.590 NABIDIA 1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.668 76.590 NABIDIA 1001284 253.434 60.199 204.903 48									
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NAMMA1001222 5.585									
NAMMA1001233 94.809 29.294 42.345 15.601 20.861 20.316 32.446 15.726	MASMA 1001220								
HAMMA1001232 130, 199 45, 692 227, 125 47, 671 38, 837 45, 692 59, 906 32, 862 MAMMA1001234 129, 344 27, 935 227, 692 95, 815 64, 344 61, 799 49, 210 34, 673 48, 484 48, 485 48, 485 48, 47 48, 485 48, 47 48, 485 48, 47 48, 485 48, 47 48, 485 48,									
NAMMA1001234 129.344 27.935 227.692 95.815 64.344 61.799 49.210 34.673 NAMMA1001237 29.560 11.083 23.224 7.241 4.489 20.199 16.883 11.003 NAMMA1001243 20.832 11.598 47.127 7.253 32.689 20.073 7.954 6.544 NAMMA1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 NAMMA1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 NAMMA1001256 169.303 81.917 266.686 187.649 131.656 44.850 55.325 59.786 NAMMA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 NAMMA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 NAMMA1001250 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 NAMMA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 NAMMA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 NAMMA1001274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 NAMMA1001280 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 NAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 NAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 NAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 NAMMA1001285 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 NAMMA1001285 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 NAMMA1001282 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001282 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378		94.809	29.294		15.601	20.861			
NAMMA1001234 129.344 27.935 227.692 95.815 64.344 61.799 49.210 34.673 NAMMA1001237 29.560 11.083 23.224 7.241 4.489 20.199 16.883 11.003 NAMMA1001243 20.832 11.598 47.127 7.253 32.689 20.073 7.954 6.544 NAMMA1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 NAMMA1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 NAMMA1001256 169.303 81.917 266.686 187.649 131.656 44.850 55.325 59.786 NAMMA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 NAMMA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 NAMMA1001250 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 NAMMA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 NAMMA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 NAMMA1001274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 NAMMA1001280 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 NAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 NAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 NAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 NAMMA1001285 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 NAMMA1001285 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 NAMMA1001282 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001282 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 NAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378	MANNA1001232	130, 199	45.692	227. 125	47.671	38.837	45.692	59.906	32.862
MANNA1001237 29.560 11.083 23.224 7.241 4.489 20.199 16.883 11.003 11.003 11.003 12.003									34, 673
MANNA1001243 20.832 11.598 47.127 7.253 32.689 20.073 7.954 6.544 MANNA1001244 44.925 10.751 11.473 9.770 11.102 14.902 16.779 4.470 MANNA1001249 43.758 23.671 15.616 19.023 10.556 26.846 10.975 13.758 MANNA1001256 169.303 81.917 266.686 187.649 131.656 44.850 55.325 59.786 MANNA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 MANNA1001260 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 MANNA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MANNA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 MANNA1001271 305.116 66.364 106.518 32.761 65.392 128.314 130.796 39.913 MANNA1001280 65.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANNA1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MANNA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.668 76.590 MANNA1001289 169.737 90.053 62.200 32.142 102.670 56.398 54.913 47.082 MANNA1001289 169.737 90.053 62.200 32.142 102.670 56.398 54.913 47.082 MANNA1001289 169.737 90.053 62.200 32.142 102.670 56.398 54.913 47.082 MANNA1001282 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 MANNA1001292 103.898 20.400 28.796 15.498 31.006 29.									
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HAMMA1001259 70.213 24.036 18.445 18.447 25.202 45.289 34.303 19.546 HAMMA1001260 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 HAMMA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 HAMMA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 MAMMA1001271 305.116 66.364 106.518 32.761 65.392 128.314 130.796 39.913 130.01274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 130.449 130.44									
HAMMA1001260 154.426 64.153 81.115 52.438 46.566 80.874 64.937 87.761 HAMMA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MAMMA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 MAMMA1001271 305.116 66.364 106.518 32.761 65.392 128.314 130.796 39.913 MAMMA1001274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 MAMMA1001280 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MAMMA1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 MAMMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MAMMA1001289 169.737 90.053 62.200 32.142 102.670 56.398 64.913 47.082 MAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 31.970 31.006 31	MAMMA 1001256	169.303	81.917	266.686	187.649	131.656	44.850	55. 325	59. 786
MANMA1001260 154, 426 64, 153 81, 115 52, 438 46, 566 80, 874 64, 937 87, 761	MAMMA1001259	70.213	24.036	18, 445	18.447	25. 202	45. 289	34. 303	19. 546
MAIMA1001262 153.326 53.618 54.054 40.354 54.252 66.416 134.449 25.835 MAIMA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 MAIMA1001271 305.116 66.364 106.518 32.761 65.392 128.314 130.796 39.913 MAIMA1001274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 MAIMA1001280 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MAIMA1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MAIMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 MAIMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MAIMA1001289 169.737 90.053 62.200 32.142 102.670 56.398 64.913 47.082 MAIMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 31.970 31.006								64. 937	87.761
MAIMA1001268 97.760 53.599 146.494 47.068 42.826 34.360 20.976 28.286 MAIMA1001271 305.116 66.364 106.518 32.761 65.392 128.314 130.796 39.913 MAIMA1001274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 MAIMA1001280 65.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MAIMA1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MAIMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 MAIMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MAIMA1001289 169.737 90.053 62.200 32.142 102.670 56.398 54.913 47.082 MAIMA1001282 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970 31.970 31.006									
MANMATOD1271 305.116 66.364 106.518 32.761 65.392 128.314 130.796 39.913 MANMATOD1274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 MANMATOD1280 65.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANMATOD1283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MANMATOD1284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 MANMATOD1286 86.284 38.290 49.421 32.175 40.490 57.656 59.470 32.210 MANMATOD1289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 MANMATOD1292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970									
HAMMA1001274 73.329 94.857 235.488 85.814 64.385 71.860 51.097 62.114 HAMMA1001280 66.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 HAMMA1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 HAMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 HAMMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 HAMMA1001289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 HAMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970									
MANMATO01280 65.399 17.595 13.218 9.853 3.831 37.015 12.303 6.374 MANMATO01283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MANMATO01284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 MANMATO01286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MANMATO01289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 MANMATO01292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970									
#AMMA1001283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 #AMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 #AMMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 #AMMA1001289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 #AMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970									
MANMATO01283 145.535 67.060 129.301 56.055 38.490 56.397 52.661 34.076 MANMAT001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 MANMAT001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MANMAT001289 169.737 90.053 62.200 32.142 102.670 56.398 64.913 47.082 MANMAT001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970	MAMMA1001280	66.399	17. 595	13.218					
#AMMA1001284 253.434 60.199 204.903 48.739 63.272 100.485 93.658 76.590 #AMMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 #AMMA1001289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 #AMMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970				129. 301	56.055	38.490	56, 397	52.661	34.076
MANMA1001286 86.284 38.290 49.421 32.175 40.490 57.666 59.470 32.210 MANMA1001289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 MANMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970									
MANMA1001289 169.737 90.053 62.200 32.142 102.670 66.398 64.913 47.082 MANMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970									
MANMA1001292 103.898 20.400 28.796 15.498 31.006 29.378 26.545 31.970	MANAGE TOU 280								
MAMMA1001296 225.022 173.717 324.251 133.662 60.125 88.173 70.926 89.316	MAMMA1001292	103.898							
	MAMMA1001296	225. 022	173.717	324. 251	133.662	60, 125	88.173	70.926	89. 316
		1 455							

Table 55

									1
	MAMMA1001298	80.876	60.189	230.669	38, 485	32.838	36.675	27.032	27. 836
	MAMMA1001305	153.258	67.563	147. 529	36. 286	31.766	65. 281	41.627	30. 730
	MAMMA1001309		8. 306	6.534	3. 627	4. 269	0.000	5, 861	6. 705
5		6.490							
	MAMMA1001310	148. 253	53.093	165.786	46.753	41.171	63. 488	82.639	54. 927
	MAMMA1001322	20.005	14.809	29.403	19.332	11. 227	14, 549	14. 163	15.700
	MAMMA1001324	82.605	28.652	85. 996	52, 506	31.339	47, 688	30.365	20.779
							97.891	81.121	27. 980
	MAMMA1001330	180.949	117.040	245. 119	52. 680	15. 121			
	MAMMA 100 1333	101.707	75. 972	213. 812	59, 950	49.965	59.640	32.340	37. 307
40	MAMMA1001334	156.564	108.340	81.315	64. 901	34.949	73.570	65. 555	73. 287
10	MAMMA1001337	105.507	35, 111	33.563	17, 119	20. 426	44.148	21.930	33.068
	MAXMA1001341	100.751	32.100	79.257	23.788	38.019	38.614	42. 286	29. 671
	MAMMA1001343	128.875	95. 425	301.822	74. 316	77. 337	85. 437	18.963	98. 899
	MAMMA1001344	32.880	35. 930	40.648	21. 963	23.320	30.315	16. 394	27. 074
	MAMMA1001345	49.749	17. 537	51.635	21, 147	20, 480	22.107	26, 805	24. 306
	MAMMA1001383	202. 565	186, 453	597. 532	117.676	100. 238	103.083	68. 993	76. 274
15									
	MAMMA1001388	149. 105	66. 100	213.624	45. 488	52.686	66.868	85. 346	57. 974
	MAMMA1001396	197.435	81.919	430. 433	80.848	94.812	95. 399	75. 293	90. 889
	MANMA1001397	116. 167	86.809	175, 125	67. 323	58.676	56. 833	61.558	52. 233
	MAMMA1001401	101.761	72.090	194. 999	62.960	48. 162	57.422	73.403	78. 023
	MAMMA1001408	62.875	17.757	62.603	9.779	13.557	44. 301	11.008	20. 408
00	MAMMA1001411	271. 344	54. 507	67.489	20.558	68.557	157. 085	134. 884	38. 338
20	MAMMA1001414	74.836	21.511	88.459	27.219	20.603	32. 791	16.798	25. 126
•	MAMMA1001415	207.635	38. 228	51.690	26.716	68.700	89. 184	99. 527	41.848
	MAMMA1001418	103.090	36. 102	91.976	39. 234	28.949	27.016	31.339	23, 195
	MAMMATCO1419	106, 299	52. 357	210.943	52.570	45. 256	41. 351	37.624	25. 914
								26. 323	
	MAMMA1001420	133.835	25. 587	149. 981	15.816	19.703	28.670		15.896
	MAMMA1001426	265. 539	180.062	165. 308	87. 320	89.096	170.869	109. 848	84.772
25	MAMMA1001428	310.313	180. 134	229. 960	136.337	147.398	262. 499	135. 345	83. 047
	MAMMA 1001432	266.375	107.317	387, 676	86.786	60.159	83.974	37. 205	60.775
	MAMMA1001435	99. 596	48, 079	193. 151	53.623	27.154	41.869	30, 388	39.835
	MAMMA1001442	103.071	100. 872	193. 544	78.030	54.054	54. 359	43.154	50.728
	MAMMA1001446					72.694	46. 485	39.641	61.589
		180.367	105, 551	197.748	98.484				
	MAMMA1001450	67.785	51.961	68.660	34. 362	32.591	32.211	28. 904	9. 424
30	MAMMA1001452	180.732	124. 244	432. 438	115.549	111.829	104. 153	92.517	96.081
• •	MAMMA1001465	528. 588	255. 549	770. 820	359. 206	364.762	388. 404	209. 219	264. 053
	MAMMA1001476	33, 639	19.551	25. 289	5.909	17. 988	24.584	26, 252	17. 981
	MAMMA1001478	117. 183	61.333	147, 393	46.785	39.649	32, 143	33,776	40.723
	MAMMA1001479	156. 131	59. 931	31.646	28.808	44.671	62.901	69. 911	26.759
	MAMMA1001487	67.613	53.042	92.480	34. 978	30.928	40.427	27.489	11.238
35									
33	MAMMA1001498	96. 522	111.213	222.159	50.813	14.811	23. 385	56, 209	28. 054
	MAMMA1001498 MAMMA1001501	96. 522 216. 969	111.213 55.879	84. 459	38. 369	49.731	88.169		28. 054 32. 036
	MAMMA 100 150 1	216.969	55.879	84. 459	38. 369	49.731	88.169	56. 209 43. 395	32.036
	MAMMA1001501 MAMMA1001502	216. 969 124. 674	55. 879 57. 815	84. 459 131. 281	38. 369 46. 452	49.731 43.478	88. 169 54. 854	56. 209 43. 395 34. 762	32.036 36.860
	MAMMA1001501 MAMMA1001502 MAMMA1001510	216. 969 124. 674 27. 993	55.879 57.815 7.591	84. 459 131. 281 13. 577	38. 369 46. 452 10. 197	49.731 43.478 11.745	88.169 54.854 6.993	56. 209 43. 395 34. 762 14. 922	32.036 36.860 8.048
	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522	216. 969 124. 674 27. 993 56. 601	55. 879 57. 815 7. 591 24. 819	84. 459 131. 281 13. 577 109. 236	38. 369 46. 452 10. 197 27. 569	49.731 43.478 11.745 21.472	88. 169 54. 854 6. 993 26. 994	56. 209 43. 395 34. 762 14. 922 29. 481	32.036 36.860 8.048 17.416
	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001529	216. 969 124. 674 27. 993 56. 601 83. 190	55. 879 57. 815 7. 591 24. 819 23. 330	84. 459 131. 281 13. 577 109. 236 52. 489	38. 369 46. 452 10. 197 27. 569 20. 883	49.731 43.478 11.745 21.472 31.879	88.169 54.854 6.993 26.994 41.170	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923	32.036 36.860 8.048 17.416 20.596
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001522 MAMMA1001529 MAMMA1001532	216.969 124.674 27.993 56.601 83.190 47.058	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881	49.731 43.478 11.745 21.472 31.879 17.641	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583	32.036 36.860 8.048 17.416 20.596 30.896
40	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001529	216. 969 124. 674 27. 993 56. 601 83. 190	55. 879 57. 815 7. 591 24. 819 23. 330	84. 459 131. 281 13. 577 109. 236 52. 489	38. 369 46. 452 10. 197 27. 569 20. 883	49.731 43.478 11.745 21.472 31.879	88.169 54.854 6.993 26.994 41.170	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923	32.036 36.860 8.048 17.416 20.596
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001522 MAMMA1001529 MAMMA1001532 MAMMA1001533	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583	32.036 36.860 8.048 17.416 20.596 30.896 16.233
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001522 MAMMA1001529 MAMMA1001533 MAMMA1001533 RAMMA1001533	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000	38.369 46.452 10.197 27.569 20.883 33.881 22.218 0.000	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001529 MAMMA1001529 MAMMA1001533 MAMMA1001533 MAMMA1001533 MAMMA1001533	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001522 MAMMA1001529 MAMMA1001532 MAMMA1001533 MAMMA1001534 MAMMA1001534 MAMMA1001534	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842	84, 459 131, 281 13, 577 109, 236 52, 489 98, 780 30, 146 0,000 23, 902 186, 325	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519	49.731 43.478 11.745 21.472 31.879 17.641 20.573 0.608 14.317 46.073	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001529 MAMMA1001529 MAMMA1001533 MAMMA1001533 MAMMA1001533 MAMMA1001533	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537
	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001522 MAMMA1001529 MAMMA1001532 MAMMA1001533 MAMMA1001534 MAMMA1001534 MAMMA1001534	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842	84, 459 131, 281 13, 577 109, 236 52, 489 98, 780 30, 146 0,000 23, 902 186, 325	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519	49.731 43.478 11.745 21.472 31.879 17.641 20.573 0.608 14.317 46.073	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660
	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001534 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001557	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781
40	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001534 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001557	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156
40	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001521 MAMMA1001522 MAMMA1001532 MAMMA1001533 MAMMA1001534 MAMMA1001534 MAMMA1001535 MAMMA1001551 MAMMA1001551 MAMMA1001551 MAMMA1001557	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 375 31. 981 39. 259	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 50. 896 115. 580	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636
40	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001532 MAMMA 1001533 MAMMA 1001534 MAMMA 1001534 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001575 MAMMA 1001575 MAMMA 1001576	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305
40	MAMMA1001501 MAMMA1001502 MAMMA1001510 MAMMA1001529 MAMMA1001529 MAMMA1001529 MAMMA1001533 MAMMA1001534 MAMMA1001534 MAMMA1001535 MAMMA1001551 MAMMA1001551 MAMMA1001551 MAMMA1001556 MAMMA1001576 MAMMA1001576 MAMMA1001576	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526 2. 133	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371
40	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001556 MAMMA 1001576 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584	216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000 214. 250	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526 2. 133 84. 714	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 245 1. 210 45. 244	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869
40	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001556 MAMMA 1001576 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526 2. 133 84. 714	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001569 MAMMA 1001576 MAMMA 1001576 MAMMA 1001584 MAMMA 1001589	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 375 31. 981 39. 259 23. 526 2. 133 84. 714	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 37. 913 27. 880	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869
40	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001532 MAMMA 1001533 MAMMA 1001534 MAMMA 1001534 MAMMA 1001535 MAMMA 1001557 MAMMA 1001575 MAMMA 1001575 MAMMA 1001576 MAMMA 1001586 MAMMA 1001589 MAMMA 1001589	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 75. 842 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 60. 438 60. 438 49. 324	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 37. 913 27. 880 35. 258	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001532 MAMMA 1001533 MAMMA 1001534 MAMMA 1001534 MAMMA 1001534 MAMMA 1001547 MAMMA 1001569 MAMMA 1001576 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001586 MAMMA 1001586 MAMMA 1001586 MAMMA 1001580 MAMMA 1001580 MAMMA 1001590 MAMMA 1001590 MAMMA 1001590 MAMMA 1001590 MAMMA 1001590	216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 29. 889 32. 647 34. 765	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016 13. 148 52. 861	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119 20.344 16.253
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001532 MAMMA 1001533 MAMMA 1001534 MAMMA 1001534 MAMMA 1001534 MAMMA 1001534 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001575 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584 MAMMA 1001584 MAMMA 1001586 MAMMA 1001586 MAMMA 1001599 MAMMA 1001606	216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 50. 539 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016 13. 148 52. 861 91. 848	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599 15. 339 88. 514	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119 20.344 16.253 78.377
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001586 MAMMA 1001586 MAMMA 1001586 MAMMA 1001589 MAMMA 1001600 MAMMA 1001600 MAMMA 1001600 MAMMA 1001606	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088 64. 637	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469 23. 619	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 49. 324 49. 324 53. 275 248. 919 74. 281	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 375 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016 13. 148 52. 861 91. 848 18. 302	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599 15. 339 88. 514 9. 100	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119 20.344 16.253 78.377
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001532 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001575 MAMMA 1001576 MAMMA 1001576 MAMMA 1001576 MAMMA 1001576 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001586 MAMMA 1001586 MAMMA 1001586 MAMMA 1001589 MAMMA 1001606 MAMMA 1001606 MAMMA 1001606	216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088 64. 637 74. 839	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 49. 324 49. 324 53. 275 248. 919 74. 281	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016 13. 148 52. 861 91. 848	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063 18. 036	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599 15. 339 88. 514 9. 100 35. 992	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192 19. 011 21. 716	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119 20.344 16.253 78.377 13.860 14.483
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001532 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001575 MAMMA 1001576 MAMMA 1001576 MAMMA 1001576 MAMMA 1001576 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001586 MAMMA 1001586 MAMMA 1001586 MAMMA 1001589 MAMMA 1001606 MAMMA 1001606 MAMMA 1001606	216. 969 124. 674 27. 993 56. 601 83. 190 47. 058 97. 390 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088 64. 637	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 765 99. 469 23. 619	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919 74. 281 9. 202	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 375 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016 13. 148 52. 861 91. 848 18. 302	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599 15. 339 88. 514 9. 100	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192	32.036 36.860 8.048 17.416 20.596 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119 20.344 16.253 78.377
40 45	MAMMA 1001501 MAMMA 1001502 MAMMA 1001510 MAMMA 1001510 MAMMA 1001522 MAMMA 1001522 MAMMA 1001533 MAMMA 1001533 MAMMA 1001534 MAMMA 1001535 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001551 MAMMA 1001576 MAMMA 1001584 MAMMA 1001584 MAMMA 1001586 MAMMA 1001586 MAMMA 1001586 MAMMA 1001589 MAMMA 1001600 MAMMA 1001600 MAMMA 1001600 MAMMA 1001606	216. 969 124. 674 27. 993 56. 601 83. 190 0. 341 32. 482 122. 717 103. 124 47. 916 137. 304 355. 571 59. 860 6. 157 150. 616 40. 717 109. 112 153. 185 217. 088 64. 637 74. 839	55. 879 57. 815 7. 591 24. 819 23. 330 33. 575 40. 032 0. 000 21. 042 75. 842 52. 282 19. 726 30. 090 57. 322 30. 398 32. 887 76. 439 29. 889 32. 647 34. 675 99. 469 23. 619 29. 828	84. 459 131. 281 13. 577 109. 236 52. 489 98. 780 30. 146 0. 000 23. 902 186. 325 155. 615 56. 549 87. 851 60. 438 0. 000 214. 250 37. 283 49. 324 63. 275 248. 919 74. 281 9. 202	38. 369 46. 452 10. 197 27. 569 20. 883 33. 881 22. 218 0. 000 24. 788 45. 519 43. 540 24. 376 31. 981 39. 259 23. 526 2. 133 84. 714 14. 016 13. 148 52. 861 91. 848 18. 302 11. 550	49. 731 43. 478 11. 745 21. 472 31. 879 17. 641 20. 573 0. 608 14. 317 46. 073 38. 692 18. 319 29. 095 62. 142 24. 246 1. 210 45. 244 19. 295 24. 411 9. 643 90. 788 10. 063 18. 036	88. 169 54. 854 6. 993 26. 994 41. 170 23. 522 25. 298 6. 274 27. 839 43. 338 47. 685 34. 666 50. 896 115. 580 30. 161 6. 758 67. 639 24. 401 44. 599 15. 339 88. 514 9. 100 35. 992	56. 209 43. 395 34. 762 14. 922 29. 481 29. 923 25. 583 46. 390 0. 000 5. 277 36. 590 20. 767 36. 128 55. 992 85. 589 16. 694 2. 949 37. 913 27. 880 35. 258 24. 456 79. 192 19. 011 21. 716	32.036 36.860 8.048 17.416 20.596 30.896 16.233 0.000 10.537 24.660 32.781 11.381 33.156 39.636 22.305 4.371 52.869 19.119 20.344 16.253 78.377 13.860 14.483

Table 56

ſ	MAMMA1001619	361.714	66.104	138, 945	35. 137	88.004	177.280	155. 721	44.365
	MAMMA1001620	113. 233	68.799	320.014	88. 182	65.387	62.891	47.797	49.428
	MAMMA1001623	32.719	16.493	22.246	8. 396	13.561	16.233	7.490	7.940
[MAMMA1001526	75. 279	8.514	13.728	10.774	12.665	55.613	57. 493	6.962
1	MAMMA1001627	28.468	7.652	39.356	8.734	4.064	8.190	14, 443	7.576
	MAMMA1001630		36, 649	115. 287	20. 971	7.371	8.511	10, 371	16.570
		35.419							
	MAMMA1001633	77.945	25. 597	143.786	22.273	51.279	40.689	37. 952	19.350
	MAMMA1001634	132.937	95. 570	297.140	83. 974	56.835	62.263	58. 952	66.333
	MAMMA1001635	140.754	47.359	225. 161	34. 126	24, 717	38.086	34, 792	34.698
								12.550	
	MAMMA1001649	30.569	12.321	20. 513	11.727	13.713	19. 299		9. 106
	MAMMA 100 16 54	150. 282	91.691	90.096	34. 969	64. 959	66.853	62.712	58. 197
	MAMMA1001660	133.470	97.805	42.199	61,020	54.089	65.813	66.019	54.874
	MAMMA1001663	394.964			154. 372	162.177	148.843	118. 542	79.262
			202. 523						
	MAMMA 100 1670	109.171	38.230	119.077	31. 362	18.030	43.797	53. 194	28. 426
	MAMMA1001671	145.809	21.188	31.621	20. 983	11.973	13.009	10,867	8.816
5	MAMMA1001679	74,490	17.313	20, 426	10.837	8. 375	23. 180	9. 271	18.786
	MAMMA 100 1683		87.078	260.375	71.605	39.630	48. 331	49.633	41.012
		147.044							
	MAMMA 1001686	12.824	14. 464	46.223	12.860	21.575	12. 528	5. 274	9.906
	MAMMA 1001688	290.960	584. 756 1	484. 182	407.762	105.060	319.616		1824.687
	MAMMA1001689	74. 586	28. 294	39.725	20.248	8. 261	19.721	31.387	18. 923
	MAMMA1001692	90.375	64.474	198.053	56. 976	35, 470	19.914	16.899	28.825
,									
,	MAMMA 1001711	111.425	82. 300	189. 195	30. 269	36.663	51. 227	10.898	27.229
	MAMMA1001715	67. 545	40. 330	71.553	28.616	19. 372	25.019	24. 223	13.907
	MAMMA 1001730	33.925	17.096	21.837	11.464	4. 477	36.743	11. 375	8. 587
	MAMMA1001735	79.384	42.172	38, 240	23.675	25. 390	20, 932	27. 963	11.313
	MAMMA1001740				17.836	17.794	23. 366	21.945	16.107
		100.894	25. 218	94. 454				21.343	
	MAMMA1001743	199.112	118.364	141.535	72.049	46. 384	86.104	96.828	100.038
5	NAMMA1001744	23.256	20. 454	0.000	2.086	2. 551	2.098	5.703	0.000
	MAMMA 1001745	121.679	94, 047	301.292	106.455	100.677	125.697	46.388	55. 894
	MAMMA1001751	58.670	37.967	90. 572	30. 921	14, 618	26.060	33.416	32.380
	MAMMA1001752	284. 221	89.024	175.680	74.746	86.008	159.864	103.908	99.685
	MAMMA1001754	57.620	30.193	53. 390	14.833	35. 182	39. 454	17.523	12.754
	MAMMA1001757	14. 456	8.290	7.632	7.247	6.076	15.580	5. 382	5. 641
	MAMMA1001760	283. 527	155. 103	596.815	118. 229	106.868	115.717	105, 154	147.707
)						5.043	11.697	22.420	16. 539
	MAMMA1001764	33.825	15.661_	33.885	14. 429				
	MAMMA1001767	41.791	27. 578	112.242	22. 484	21.848	16.357	11.576	9. 367
	MAMMA1001768	50.861	34, 645	129.707	25, 692	23.037	24.674	27.811	11.075
	MAMMA1001769	206.737	82.818	645. 195	110,913	102.640	105.607	80.653	102.144
					16.877	55.099	52.348	41.113	48.806
	MAMMA1001771	123. 973	30. 551	49.772					
	MAMMA1001773	47.743	27. 204	35, 277	8, 450	18.002	17, 141	23.713	30.755
)	MANNA1001778	104.585	49.619	92. 589	42. 249	35.085	50.584	39.215	26.862
	MAMMA1001783	140.821	89.274	371.095	82.231	85.003	87.248	61.999	71.448
	MAMMA1001785	119.072	65.819	256.400	60. 491	37.351	65, 802	45.875	54.652
					9.749	9.870	11.494	13.172	10.408
	MAMMA1001788	37. 967	8. 305	25. 708					
	MAMMA1001790	202.092	181. 258	279.482	57.700	22.737	29. 284	28.819	46.106
	MAMMA1001800	24. 282	11.444	30. 466	12.517	1.763	8.501	13.065	25.671
)	MAMMA1001804	150.744	16.771	51.213	14. 975	33.630	67.533	64.799	20.701
	MAMMA1001806	62.312	54. 896	146.142	37, 371	11.402	36.501	43.675	52.846
		17.002		32.023	10.166	5. 995	9.576	10.245	11.255
	MAMMA1001812		11.569						
	MAMMA1001815	50.743	27. 272	61.778	19.704	15, 636	25.863	15. 187	22.130
	MAMMA1001817	10.653	7.578	15. 446	7.044	7.758	3.611	7.974	11.601
	MAMMA1001818	48.733	19.657	87. 193	21.647	18.566	18.770	19.255	18.678
	MAMMA1001819	165. 340	99. 233	343, 318	111.523	112.261	57.848	73.268	87.725
)									
	MAMMA1001820	48.662	22.951	34. 879	16. 243	11.743	9, 468	15.897	11.396
	MAMMA1001824	125.683	53.824	187. 383	58. 214	53.691	47.999	45. 347	37. 548
	MAMMA1001832	56.633	30.370	42, 082	21.957	23.518	23.996	20.046	8. 482
	MAMMA1001836	128. 477	58. 280	179.541	45. 913	43, 465	44.952	56,814	24. 346
				172.658	60. 299	38.153	37.090	17.947	50.301
	MAMMA1001837	118.428	66.031						
)	MAMMA1001848	42. 562	27. 622	82.759	24, 693	20.435	22.941	15, 102	19.124
	MAMMA1001850	402.506	243. 182	312.586	171. 182	143.034	232.615	91, 466	106.637
	MAMMA1001851	123. 305	30.035	69.870	64.763	41.560	39, 454	33, 329	45. 924
	MANNA 100 1031			321.896		133.655	112.820	91.724	115. 602
	MAMMA1001852	198.774	161.311					39.634	
			1 137 469			77. 240	070	AFA UF :	45. 321
	MAMMA1001854	158.894	117.462	234. 984			42.929		
	MAMMA1001854 MAMMA1001858	148. 310	133.834	240. 344		24.063	35.871	73. 151	58. 279

Table 57

IAMMA1001864	169,742	52.389	185, 785	37. 880	50.896	67.999	55. 272	23, 142
AMMA1001868	82.643	56.439	59. 491	62.418	34. 438	47.003	29.588	35. 585
					11.275		7. 189	10.453
AMMA1001874	9.192	9.651	51.178	7.405		9.054		
AMMA1001878	190.515	70. 315	227.600	164.835	101.886	72.219	79.645	146. 982
188 (001 AMMA)	159.918	94. 489	292.528	95. 467	48. 528	98. 588	39.271	81. 114
IAMNA1001885	117.729	44. 975	110.656	53. 460	26.142	52. 223	41.423	29.156
IAMMA1001890	127. 969	47.712	247. 654	60. 558	29. 367	36.838	39, 109	41.483
						30.030		
AAMA 1001893	90. 120	22. 271	50. 435	19.070	23. 222	27.783	36.643	18.711
(AMMA1001901	78.854	67.274	188.894	57.356	38. 856	45.633	22.050	26.367
MANNA 1001907	159.767	70.062	305. 846	76.004	91.563	25.690	68. 288	28. 595
MAMMA1001908	44.964	27. 928	41. 967	55.852	40.219	53.008	32, 123	40.375
			10 100			0.000	0.000	5. 175
MAMMA1001919	0.000	82.865	12.109	0.000	2.270			
MAMMA1001931	59.705	9.869	29. 213	49, 582	13.981	18. 165	29. 466	11.467
MANIMA 1001937	47.045	26.453	33. 302	16.535	17.844	31.265	29.899	19.650
MAMMA1001951	114.033	76.574	311.618	70. 531	55, 661	40.552	39.990	40. 224
MAMMA1001956	171.199	78.116	295.630	76. 171	65, 654	47.426	67.568	57.411
			155.366	46.819	41.429	43. 671	26. 153	26. 982
MAMMA1001957	114.304	40.789						
MAMMA1001960	99.822	53.449	192.955	55. 422	57. 938	23. 395	42.027	44. 844
MAMMA1001963	6.938	3.651,	9.748	3.671	3. 337	0.000	0.000	5. 275
MANNA 1001969	237.109	164.919	517.768	178. 594	149.500	109. 284	97.612	137, 120
MAMMA1001970	199.358	123.085		101.158	41.691	71.806	71.685	61.125
					1.081	1. 561	0.000	0,000
MAMMA1001978	1.206	0.000	0.000	0.000				
MAMMA1001992	189. 502	91.630	283.440	78.807	70.640	63.218	71. 282	32.898
MAMMA1001994	85. 231	21.385	143, 259	40.178	38. 484	54. 686	24.893	33.837
MANNA 1002008	66.834	77.793	37.547	14.813	20.016	33. 334	39.365	10.388
MAMMA1002009	144. 462	65.030		107.350	55.438	47. 107	40, 434	57.138
			27.624		19.701	17.344	22. 354	14.449
MAMMA1002011	32.832	13. 901		10. 188	19.101			
MAMMA1002022	107.727	67.057	159.576	65. 640	59. 239	37. 381	36, 122	50.747
MAMMA1002024	176.885	70, 125	207. 390	72.614	55.279	78. 953	108, 945	46.948
MAMMA1002032	270. 523	130.983	362.313	98.620	95. 826	104.970	73.966	83.780
MAMMA1002033	132.652	119.984	303.660	81.264	93.758	74. 391	34, 919	49.831
		15.313	18, 901	14.070	10.859	15, 705	11.098	10.476
MAMMA1002041	19.611							
MAMMA1002042	78.700	42, 958	161.397	37.566	30. 208	55. 486	24.562	23.890
MAMMA1002645	7.131	8.948	24.018	14. 459	14.811	11.172	1, 533	10.371
MAMMA1002047	82.875	57.343	192.240	55.806	45. 781	34.315	27.824	37.210
MAMMA1002056	212. 189	152. 323	474.785	146.238	94.617	84.218	104.806	75.923
MAMMA1002058	149.112	126.148	334.116	98. 541	74.809	81.670	44. 227	65. 825
					5.782	5. 917	16. 902	5. 536
MAMMA1002060	13.278	7.931	14, 514	12.643				
MAMMA1002065	128. 185	46.405	127.810	82.855	59. 107	72.737	63.052	39.667
MAMMA1002068	110.652	64. 982	163.753	51. 583	45. 893	40.656	37.400	24. 128
MAUMA1002070	61.186	24.791	29. 988	16.102	15.306	31.362	22.002	21. 338
MAMMA1002078	170. 197	38.633	93.014	30.633	33.682	90.533	42.110	14. 299
				10. 208	14.094	14.792	10.377	10.263
MAMMA 1002080	21. 195	14. 596	12.645					
MAMMA 1002082	111.870	77.716	117.819	55.009	54. 940	28. 457	25. 946	21. 254
MAMMA1002084	74. 297	40.086	152.790	30.118	30.052	28. 788	24. 428	24.140
MAMMA1002087	17. 991	17.619	30.479	8.932	13.026	13. 365	9, 996	6.344
MAMMA1002091	78.604	26.611	41.258	17.086	26.812	39.757	46.803	27.660
MAMMA1002093	17. 498	0.000	5. 942	5. 592	5.630	8. 103	11.278	4.689
MAMMA1002095		13,430	22.728	13.058	20.650	32. 157	32.621	8. 152
	78.790						27.923	11.735
MAMMA1002108	91.919	6.035	31.027	13.639	7.939	32.486		
MAMMA1002112	24.376	27.337	10.667	11.574	5. 250	15.678	14.329	37.463
MAMMA1002118	12.060	5.100	8.756	5. 943	6. 502	7.856	7.396	3.149
MAMMA1002119	122.271	36.908	59.513		36.895	38, 172	39.046	32.476
MAMMA1002125	159. 277	83.844	373.786		54. 991	63.367	35, 366	35.797
			431.047		117.027	84.728	70. 558	62. 381
MAMMA1002126	231.380	139. 298					39, 193	
MANMA1002128	102.647	35.864	48. 863		20.911	44. 235		25, 406
MAMMA1002132	226.752	118.230	198.712		88.860	84.266	50.630	48. 550
	54.642	53.227	115.593	42. 121	33. 524	31.026	24. 905	32.121
MAMMA 1002140	121.646	33, 512	49, 214		27. 295	103.698	68. 348	39.850
MAMMA1002140	1 161.070			38.118	5. 895	13.974	10.806	45. 937
MAMMA1002142		1 15 750		0. 110	1 3.033	1 13.314	1 10.000	
MAMMA1002142 MAMMA1002143	150. 595	15.368				67 676	74 24	99 493
MAMMA1002142 MAMMA1002143 MAMMA1002145	150. 595 237. 202	72. 397	165. 166	45. 537	53.986	87.872	73.605	
MAMMA1002142 MAMMA1002143	150. 595		165. 166	45. 537		53. 571	33.082	8.766
MAMMA1002142 MAMMA1002143 MAMMA1002145	150. 595 237. 202	72. 397	165. 166 45. 076	45. 537 27. 984	53. 986 33. 648			8. 766 19. 099

Table 58

MAMMA1002156	- 3 616 T	0 000 1	14.013	0.880	0.000	0.000	0.000	0.554
	3.612	2.088						
IAMMA1002158	70.916	40.655	88. 575	24. 125	14.786	21.144	21.721	31.526
AMMA1002164	109. 211	29. 584	54. 163	32.089	28.633	66.844	29. 378	23.138
ANNA 1002165	166.029	111.787	135. 468	73,710	66.970	77, 137	88, 540	53.125
IAMMA1002170	0.000	0.000	0.000	1.159	0.000	0.000	0.000	0.000
IAMMA 1002174	139, 902	178. 299	326. 262	182.252	147.225	141.399	87.695	71.650
			21.290	16.108	13.918	22.449	12.876	18.564
MAMMA1002175	49. 535	20.661	21.230		30.704			
MAMMA 1002180	117.470	55.089	69.154	18.969	36.764	45, 946	59.721	45. 237
MANMA1002198	123. 227	67. 539	235.488	54.699	51.835	48.796	31.324	62.413
		63. 437	420.688	47.331	61.775	61.499	42. 296	74.029
	114.861							
MAMMA1002206	86.539	30, 665	50.318	17.788	32. 139	63. 320	64. 272	56. 392
MAMMA1002209	124.961	73.557	143.211	32.601	43.486	64. 448	43.661	36. 987
		148. 590		150.983	162.248	310.059	210.563	225.764
	446.836					50.008	34. 183	47.670
MAMMA 1002219	103.054	68.338	110.047	29. 595	35.094			
MAMMA1002224	155, 329	135.036	325.596	92.243	139.113	54. 888	50.692	104. 338
MAMMA1002229	54.055	19.297	24.594	8, 408	18.280	19.024	14.880	18.482
					50. 164	62. 315	35. 205	65.871
	131.172	95.706	345. 936	76.632				
MAMMA 1002233	40. 299	20. 503	27.780	14.645	13.380	24. 157	18.866	16.294
MAMMA 1002234	16.951	13.815,	19.460	7. 251	4. 128	10.631	13.812	19. 438
		43 653	60 603		51.817	24. 897	29.324	44.837
MAMMA1002236	50.642	23.553	50.683	14, 162	31.011			
MAMMA1002243	88.955	30.943	38. 127	26.451	21.889	37. 268	32.369	10.849
MAMMA1002250	101.569	23.851	171.031	56.513	74.300	48. 863	11.431	66.114
MAMMA1002253	515. 165	161.871	322.750	80.630	175.660	370.878	217.429	157.156
				00.030				
MAMMA1002267	129. 167	239.800	180.046	95.357	56.654	98. 387	72.076	331.998
MAMMA1002268	36.456	15.771	39.216	17.501	24.043	16.873	20.704	13.929
MAMMA1002269	27.848	6. 625	13.419	16.093	10.154	9.666	6.915	4.635
				38.160	60.059	34.106	22.977	37.892
MAMMA1002282	53.648	58. 269	178. 298	38, 100				31.032
MAMMA1002292	62.491	17.873	48. 526	22.803	16.647	14.012	30.027	30.270
MAMMA1002293	236.280	162.513	481,000	154. 526	85. 449	104.060	60.152	54.729
		24.654	124.002	36.492	33.138	43.853	25. 143	19.816
MAMMA1002294	110.705				10 100			17.505
MAMMA1002297	66.424	40.774	88. 229	32.940	16.126	21.061	14. 524	
MAMMA1002298	104.368	30.772	64.493	24.071	29.853	40. 308	35.653	29.912
MAMMA1002299	102.764	41.185	67. 139	29.656	30.944	33.813	19.722	23.248
					20. 255	27.771	17. 935	16.223
MAMMA 1002308	69. 299	30.798	86.503	30.668	29.756			
MAMMA1002310	494. 257	272.509	645.571	186.568	219.463	344.867	183. 571	203.149
MAMMA1002311	151.653	60.941	315.707	69.190	66.700	63.609	50. 563	40.723
		36.483	113.839	34.110	19.878	36.852	19, 114	16.993
MAMMA1002312	79.548							20. 920
MAMMA1002317	96.094	32.026	188.632	45. 170	46.365	46. 409	41.391	
MAMMA1002319	141.320	69.599	218.472	74.218	50.463	59. 927	44. 261	42.418
MAMMA 1002322	144. 393	65, 401	253.730	67.857	46.931	25. 375	51.002	44. 820
				17 007		27 210	20. 223	13.61
MAMMA 1002329	49.002		28. 349		21.239	27. 218		
MAMMA 1002332	55.840	30.915	137.766	47.492	35. 312	32. 956	23, 130	16.41
MAMMA 1002333	75.478	17.882	32.309	19.280	28. 575	31, 145	41.629	17.63
		50.373	149.587	54.778	40. 367	18.695		26.410
MAMMA1002335	171.866				1 70. 301	46 025		
MAMMA 1002339	91.741	62.618	152.049	63.915	53.097	48.035	33.591	31.79
MAMMA 1002347	98.915	55, 800	120.784		55. 929	33.327	45. 235	27.50
MAMMA1002351	70.045	22.016	35, 600		20. 122	33.583	21.722	19.63
MANATA I VVATA		17.786	22.690	23.069	12.412	24.411	13.818	11.94
MAMMA 1002352	52. 143	11.100			40 50	100 000	12, 132	34.57
MAMMA 1002353	128. 336	52.785	144.030	46. 481	46.561	36.806		
MAMMA 1002355	46. 995	34.505	123.684	29.737	22.025	29. 352		22.66
MAMMA 1002356	40. 901	21.732	86. 932		25. 451	22.826	13, 215	18.95
					142.084	59.794	89.656	
MAMMA1002359	276.825	92. 529	330.418					10 10
MANMA 1002360	42.725	25.740	47. 382	16.661	18. 409	9. 982	9. 481	12.12
INCOME LIBERTY I VOLUME	152.118	88.131	201.317		41.767	51.778	26.886	27.24
					14.517	23, 579	14.318	19.59
MAMMA1002351		72 602		1 41.134		65, 479	60. 201	
MAMMA1002361 MAMMA1002362	39. 281	22.692						210.78
MAMMA1002351		22.692 75.867	50.909	48. 285	31.065			
MANMA1002361 MANMA1002362 MANMA1002367	39. 281 142. 262	75.867	50.909	48. 285		49. 599	32.494	49.25
MANMA1002361 MANMA1002362 MANMA1002367 MANMA1002371	39. 281 142. 262 119. 755	75.867 66.644	50.909 278.090	48. 285 138. 658	42.317	49. 599	32.494	
MAMMAT 002361 MAMMAT 002362 MAMMAT 002367 MAMMAT 002371 MAMMAT 002380	39. 281 142. 262 119. 755 90. 587	75.867 66.644 47.691	50.909 278.090 161.106	48. 285 138. 658 38. 559	42.317 31.139	49. 599 36. 350	32. 494 34. 696	25. 22
MANMA1002361 MANMA1002362 MANMA1002367 MANMA1002371	39. 281 142. 262 119. 755	75.867 66.644	50.909 278.090 161.106	48. 285 138. 658 38. 559 71. 113	42.317 31.139 46.508	49. 599 36. 350 40. 126	32. 494 34. 696 29. 975	25. 22 44. 41
MAMMA1002361 MAMMA1002362 MAMMA1002367 MAMMA1002371 MAMMA1002380 MAMMA1002384	39. 281 142. 262 119. 755 90. 587 90. 935	75.867 66.644 47.691 85.538	50. 909 278. 090 161. 106 249. 278	48. 285 138. 658 38. 559 71. 113	42.317 31.139	49. 599 36. 350	32. 494 34. 696	25. 22 44. 41
MAMMA 1002361 MAMMA 1002362 MAMMA 1002367 MAMMA 1002371 MAMMA 1002380 MAMMA 1002384 MAMMA 1002385	39. 281 142. 262 119. 755 90. 587 90. 935 13. 712	75.867 66.644 47.691 85.538 7.306	50. 909 278. 090 161. 106 249. 278 6. 051	48. 285 138. 658 38. 559 71. 113 7. 420	42.317 31.139 46.508 3.720	49. 599 36. 350 40. 126 9. 599	32. 494 34. 696 29. 975 8. 116	25. 22 44. 41 7. 60
MAMMA1002361 MAMMA1002362 MAMMA1002367 MAMMA1002371 MAMMA1002384 MAMMA1002385 MAMMA1002385	39. 281 142. 262 119. 755 90. 587 90. 935 13. 712 119. 086	75.867 66.644 47.691 85.538 7.306 26.468	50. 909 278. 090 161. 106 249. 278 6. 051 66. 535	48. 285 138. 658 38. 559 71. 113 7. 420 12. 989	42.317 31.139 46.508 3.720 40.464	49. 599 36. 350 40. 126 9. 699 53. 956	32.494 34.696 29.975 8.116 37.080	25. 22 44. 41 7. 60 19. 51
MAMMA 1002361 MAMMA 1002362 MAMMA 1002367 MAMMA 1002371 MAMMA 1002380 MAMMA 1002384 MAMMA 1002385	39. 281 142. 262 119. 755 90. 587 90. 935 13. 712	75.867 66.644 47.691 85.538 7.306	50. 909 278. 090 161. 106 249. 278 6. 051 66. 535 97. 224	48. 285 138. 658 38. 559 71. 113 7. 420 12. 989 19. 547	42.317 31.139 46.508 3.720 40.464 21.438	49. 599 36. 350 40. 126 9. 699 53. 956 26. 503	32. 494 34. 696 29. 975 8. 116 37. 080 20. 868	25. 22 44. 41 7. 60 19. 51 14. 25
MAMMA1002361 MAMMA1002362 MAMMA1002367 MAMMA1002371 MAMMA1002384 MAMMA1002385 MAMMA1002385	39. 281 142. 262 119. 755 90. 587 90. 935 13. 712 119. 086	75.867 66.644 47.691 85.538 7.306 26.468	50. 909 278. 090 161. 106 249. 278 6. 051 66. 535 97. 224	48. 285 138. 658 38. 559 71. 113 7. 420 12. 989 19. 547	42.317 31.139 46.508 3.720 40.464 21.438	49. 599 36. 350 40. 126 9. 699 53. 956	32.494 34.696 29.975 8.116 37.080	49.25 25.22 44.41 7.60 19.51 14.25 53.90

Table 59

MANMA1002400	10.797	7.113	11, 587	4.041	5.847	4, 732	4.516	4. 19
MAMMA 1002409	93.810	75.886	50. 232	41.725	30.159	43, 673	520.771	70. 37
MANMA1002411	81.111	34, 713	76. 973	23. 185	26, 301	31, 997	16. 726	11.90
MAMMA1002413	199.066	68.034	377. 354	55. 454	56.059	50.318	26. 763	38. 96
MANMA1002417	30. 976	26. 195	58. 136	15. 593	17,649	14. 266	7.765	11.3
MAMMA1002427	87, 721	47.715	208. 629	48. 123	38.391	40.117	26. 156	31.5
MANNA1002428	108. 360	83.671	293. 146	88.263	84. 156	51.786	57. 518	57. 1
MANMA1002433	90. 843	23. 726	38. 263	19.586	19.565	44. 397	36. 529	25.0
MAMMA1002434	117. 152	72.024	272. 113	68.694	66.706	54. 616	45. 191	46.5
MAMMA 1002446	102.855	36. 748	90.796	22. 955	36.351	49. 598	42.676	12.8
MANMA 1002 447	77. 962	49. 457	171.445	42.653	21.446	36.510	25. 929	27. 9
MANIMA 1002 454	314. 500	201.950		188.845	118.797	99.696	72.794	103.9
MANMA1002461	204, 581	47. 899	153.652	28. 137	56.943	63.968	55. 245	48.4
MANNA 1002463	130. 489	40.148	72.561	25.745	31.969	67.395	41.920	28.7
MANONA 1002464	94. 697	34. 520	44. 484	18, 573	24.045	50.857	37. 103	17.4
MAMMA 1002466	27, 080	25. 120	36. 208	16.549	16. 920	44. 337	37.029	13.8
MAMMA1002470	66. 277	10. 542	19. 623	14.778	9. 384	20.022	21.241	15.3
MAMMA1002475	35. 982	26.009	77.707	23.670	24.685	10.963	12. 591	26.3
MAMMA1002480	85. 342	48. 419,	144. 499	40.755	50.788	48. 101	35. 187	30.0
MAMMA1002485	256, 024	56. 235	75. 461	32.978	72.095	120.038	77.311	49. 9
MAMMA 1002494	66.749	23. 381	164.418	25. 376	48. 947	43.136	11.733	14. 4
MAMMA1002498	58.032	20. 346	24. 265	12.932	13. 125	26. 950	19.794	5.5
MAMMA1002524	73.628	20.842	11.923	21.047	20.268	27.749	12.366	14.6
MAMMA 1002530	82.789	19.903	43.603	13.551	9, 151	28. 535	27.989	12.5
MAMMA1002538	101.182	27.725	28. 460	21.181	31.900	45. 529	26.380	25. 6
MAMMA 1002545	131.415	100.020	322. 993	72.173	54. 265	23.145	30.820	51.3
MAMMA1002554	51.033	30.923	62. 549	16. 548	18.644	38. 344	32.052	17.4
MAMMA1002556	201.613	62.773	211.073	70.139	99.337	37. 921	45. 357	46.5
MAMMA1002561	199.748	128, 004		135, 854	118. 280	54.740	81.217	51.6
MAMMA1002565	57. 918	43. 508	20. 564	13. 434	36, 930	27. 532	51.392	13.7
MAMMA1002566	29. 155	16. 405	7. 906	3.450	1.967	13.518	5.709	5.3
MAMMA1002571	73.034	22. 187	37. 154	25. 594	6.079	28.030	19. 946	20.9
MANMA1002573	218. 479	62.669	183. 544	61.350	46.029	113.781	65.617	60.5
MAMMA1002576	109.621	18, 498	33.802	10.617	22.615	43. 283	55. 199	26.4
MAMMA 1002584	244. 467	197. 626	384.879	79.185	103. 251	112.917	113. 914	151.6
MAMMA 1002585 MAMMA 1002586	133. 865	28. 963	56. 983	17. 186	16.306	13.727	51.687	25.7
MANMA1002589	67. 168 98. 120	39. 043 25. 567	34.776	15.656	19. 252	29. 596	35. 555	19.9
MAMMA1002590	268. 176	57.804	26.638 202.329	16.923 36.276	18.956	18. 249	16. 364	12.5
MAMMA1002593	131. 425	64. 951	130. 257	54. 131	77.487 23.515	180. 923 55. 983	123.883 37.410	42.5
MAMMA1002597	76. 091	50. 352	131.097	33.606	42.551	25. 425	36.396	36. 2 34. 7
MAMMA1002598	69. 190	45. 133	59. 324	58.225	35. 339	68.531	47. 164	70. 2
MAMMA1002603	122. 932	40. 124	155. 801	51.386	48.672	98.075	64. 732	66. 1
MAMMA1002612	330. 999	152. 583	441.574	105.603	112.764	175. 106	98. 853	99.4
MAMMA1002617	363, 139	211.631	557.754	145. 485	146. 260	203.052	110.009	118.2
MAMMA1002618	90. 423	66.208	129.807	53.454	46.096	53. 758	43. 899	55.8
MAMMA1002619	34. 076	14. 223	23. 292	10.350	14.540	15. 236	12.465	13.6
MAMMA1002622	112.756	60.308	263. 518	46.461	43.508	41.984	32.044	52.6
MAMMA1002623	89.689	68.083	149.811	64.401	102.216	102. 611	54. 682	73.3
MANMA1002625	83.660	44. 949	94.038	26.154	32.540	34. 576	38. 497	28. 1
MAMMA1002627	9, 090	2.616	7. 631	2.675	0.000	3.940	7.852	8.8
MAMMA1002629	111.050	96. 279	397. 433	77.573	45.933	89.752	53.737	108.3
MAMMA1002631	50. 470	10.960	11.524	6.679	3.741	10.219	10.741	11.3
MAMMA1002633	32. 234	20. 386	37.729	16.053	9. 358	12.456	8. 581	32.1
MAMMA1002636	59.898	50. 529	142. 123	25.014	15.348	18.150	38.018	22.6
MAMMA1002637	58. 583	21.541	11.323	5.892	14.789	18.069	26.406	22. 1
MAMMA1002646	55. 442	29.770	36. 308	23.176	15.750	18.816	26. 997	38.8
MAMMA1002648	49.661	48. 800	69. 217	43.621	54.730	39. 438	38. 742	48.0
MAMMA1002650	15. 384	6.907	9. 595	4.820	3. 958	6.140	8. 225	6.0
	61.935	69. 556	44. 994	60.882	59.089	42. 135	62.414	54.6
MAMMA1002652			1 12 (00	11.569	8.462	23.347	10.991	22. 1
MAMMA 1002655	49.617	25. 105	13. 568					
MAMMA 1002655 MAMMA 1002662	122.410	44. 430	94. 935	34.850	32.770	58.417	41.476	
MAMMA 1002655								39. 9 153. 3 40. 5

Table 60

NAMAH 1002788 169 486 32 550 56. 424 32 013 39 387 88 586 48 0. 699 45 05 NAMAH 1002885 25 207 18 401 21 785 11 31 18 28 31 402 25 560 25 200 NAMAH 1002885 27 274 9 361 3 697 10 386 2 003 4 100 3 302 8 NAMAH 1002888 39 272 32 200 43 587 33 15 54 10 571 20 167 32 429 30 79 NAMAH 1002889 38 348 10 645 5 272 3 333 31 35 35 11 1796 12 322 32 30 30 30 30 30	AMAMA 100268 2 7, 774 9.361 3, 887 86, 864 80, 699 45, 058 AMAMA 100268 2 7, 774 9.361 3, 887 10, 386 2, 003 4, 100 3, 302 9, 849 AMAMA 100268 3 18, 774 9.361 3, 887 10, 386 2, 003 4, 100 3, 302 9, 849 AMAMA 100268 3 18, 772 32, 200 43, 557 33, 153 4, 354 11, 796 12, 328 34, 789 AMAMA 100268 3 18, 772 32, 200 43, 557 33, 153 4, 354 11, 796 12, 328 34, 789 AMAMA 100268 13, 348 10, 845 5, 772 3, 313 3 2, 314 3, 828 12, 779 6, 883 AMAMA 100270 8 232, 250 119, 730 163, 346 75, 850 55, 245 76, 116 103, 624 109, 697 AMAMA 100271 128, 862 101, 841 359, 100 105, 555 79, 900 76, 543 28, 135 56, 135 14, 344 AMAMA 100271 128, 862 101, 841 359, 100 105, 555 79, 900 76, 543 28, 135 56, 135 14, 344 AMAMA 100271 128, 862 101, 841 359, 100 105, 555 79, 900 76, 543 28, 135 56, 137 57, 914 AMAMA 100271 128, 862 101, 841 359, 100 105, 555 79, 900 76, 543 28, 135 56, 137 57, 914 AMAMA 100271 128, 862 107, 741 37, 741 37, 744 27, 554 13, 186 13, 183 34, 194 740 33, 088 AMAMA 100271 128, 862 107, 800 360, 516 86, 920 49, 926 57, 925 48, 421 76, 576 AMAMA 100272 1 128, 620 78, 060 360, 516 86, 920 49, 926 57, 925 48, 421 76, 576 AMAMA 100272 1 128, 620 778, 060 360, 516 86, 920 49, 926 57, 927 41, 927 40 33, 088 AMAMA 100272 1 128, 620 778, 060 360, 516 86, 920 49, 926 57, 927 41, 927 40 33, 088 AMAMA 100272 1 128, 620 778, 060 360, 516 86, 920 49, 926 57, 927 41, 927 40 33, 088 AMAMA 100272 1 128, 620 778, 060 360, 516 86, 920 49, 926 57, 927 41, 927 40 33, 088 AMAMA 100272 1 128, 620 778, 060 360, 516 86, 920 49, 926 57, 927 41, 927 40 33, 088 AMAMA 100272 1 128, 620 778, 060 360, 516 86, 920 49, 926 57, 927 41, 927 40 33, 088 AMAMA 100272 1 128, 620 778, 620 360, 160, 160, 160, 160, 160, 160, 160, 1									
NAMAH 1002584 189, 486 32, 550 50, 424 32, 013 39, 987 88, 586 480, 699 45, 051 NAMAH 1002585 25, 020 18, 401 21, 755 11, 315, 31 11, 628 31, 402 25, 565 25, 020 NAMAH 1002585 125, 020 18, 401 31, 602 31, 400 33, 02 38, 402 NAMAH 1002585 18, 717 82, 318 15, 541 32, 934 10, 671 20, 167 72, 429 30, 79 NAMAH 1002585 18, 717 12, 23, 18, 18, 18, 18, 18, 18, 18, 18, 18, 18	AMAMA 100268 2 1, 274 9, 361 3, 687 10, 386 2, 003 4, 100 3, 302 9, 849 AMAMA 100268 2 1, 274 9, 361 3, 687 10, 386 2, 003 4, 100 3, 302 9, 849 AMAMA 100268 2 1, 274 9, 361 3, 687 10, 386 2, 003 4, 100 3, 302 9, 849 AMAMA 100268 3 32, 272 32, 200 43, 657 33, 153 4, 254 11, 795 12, 128 34, 400 AMAMA 100268 3 33, 272 32, 200 43, 657 33, 153 4, 254 11, 795 12, 128 34, 400 AMAMA 100268 3 33, 272 32, 200 43, 657 33, 153 4, 254 11, 795 12, 128 34, 400 AMAMA 100276 8 13, 250 119, 730 163, 346 75, 580 55, 245 76, 116 103, 624 109, 697 AMAMA 100271 128, 882 101, 814 359, 100 105, 555 79, 900 76, 543 25, 135 51, 955 AMAMA 1002712 55, 151 50, 304 36, 811 8, 507 18, 857 75, 978 44, 035 47, 031 AMAMA 1002711 128, 862 101, 814 359, 100 105, 555 79, 900 76, 543 25, 135 51, 978 AMAMA 1002712 128, 562 178, 060 360, 516 86, 920 49, 226 57, 925 48, 421 76, 576 AMAMA 1002721 128, 862 103, 874 118, 874 13, 366 33, 383 393 74, 740 33, 038 AMAMA 1002721 128, 862 103, 874 118, 974 118	IAMMA1002673	94. 294	135, 347	302.435	85.978	116.544	122.876	58.765	72.402
MANDALODZESS 25,020 18,401 21,785 11,312 11,628 3,402 5,660 25,000 MANDALODZESS 7,7274 9,161 3,697 10,387 10,387 10,386 2,003 4,100 3,302 5,460 MANDALODZESS 65,711 52,319 15,641 12,934 10,671 20,167 32,429 30,79 LANDALODZESS 18,571 15,2319 15,641 12,934 10,671 20,167 32,429 30,79 LANDALODZESS 18,348 10,645 5,272 3,333 2,314 3,625 12,679 6,38 MANDALODZESS 18,348 10,645 5,272 3,333 2,314 3,625 12,679 6,38 MANDALODZESS 11,1312 11,131 16,328 3,393 57,919 29,820 55,14 MANDALODZESS 11,1312 11,131 16,384 57,585 65,245 75,116 103,624 109,58 MANDALODZESS 11,1312 11,131 16,384 57,585 65,245 75,116 103,624 109,58 MANDALODZESS 13,131 50,304 3,681 18,507 18,357 75,116 103,624 109,58 MANDALODZESS 13,131 50,304 3,681 18,507 18,357 75,116 103,624 109,58 MANDALODZESS 13,131 50,304 36,811 8,507 18,357 75,578 14,085 47,08	AMAMAIODZESS 25,020 18.401 21.785 11.312 11.628 3.402 5.660 25.002 AMAMAIODZESS 67.7274 9.361 3.897 10.386 2.003 4.100 3.302 9.846 AMAMAIODZESS 67.71 52.319 15.641 32.934 10.671 20.167 32.429 30.795 AMAMAIODZESS 36.711 52.319 15.641 32.934 10.671 20.167 32.429 30.795 AMAMAIODZESS 91 18.348 10.645 5.272 3.333 2.314 3.625 12.679 6.883 AMAMAIODZESS 91 18.348 10.645 5.272 3.333 2.314 3.625 12.679 5.883 AMAMAIODZESS 91 18.348 10.645 5.272 3.333 2.314 3.625 12.679 5.883 AMAMAIODZESS 91 18.348 10.645 5.272 3.333 2.314 3.625 12.679 5.883 AMAMAIODZESS 91 19.700 163.846 75.850 55.245 76.116 103.624 105.697 AMAMAIODZESS 91 19.700 163.846 75.850 55.245 76.116 103.624 105.697 AMAMAIODZESS 91 19.700 163.861 8.507 18.857 75.978 44.065 47.001 AMAMAIODZESS 91 12.8520 17.841 37.741 37.674 23.554 13.366 139.383 49.740 33.088 AMAMAIODZESS 91 17.852 18.508 63.389 11.854 18.8507 18.857 75.978 14.065 47.001 AMAMAIODZESS 91 19.868 183.567 75.931 18.66 139.383 49.740 33.088 AMAMAIODZESS 91 19.868 183.567 75.931 18.852 77.925 14.842 11.656,771 12.8520 18.854 18.8507 18.857 79.931 10.8740 18.857 18.951 18.958 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 18.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 17.758 18.958 1	MANNA 1002684				32.013	39. 987	86.564	80.699	45, 058
MAINAI DOZESS 7, 274 9, 361 3, 697 10, 386 2, 003 4, 100 3, 302 3, 44 MAINAI DOZESS 56, 771 52, 339 15, 641 12, 934 10, 671 20, 167 21, 429 30, 791 MAINAI DOZESS 36, 771 52, 339 15, 641 12, 934 10, 671 20, 167 21, 429 30, 791 MAINAI DOZESS 39, 272 32, 200 43, 657 33, 153 4, 354 11, 796 12, 328 34, 40 MAINAI DOZESS 31, 348 10, 645 5, 272 33, 313 2, 314 3, 625 12, 679 6, 88 MAINAI DOZESS 31, 373 107, 821 376, 150 82, 188 33, 933 57, 919 29, 820 551, 44 MAINAI DOZESS 32, 250 119, 730 153, 846 75, 850 65, 245 75, 116 103, 624 109, 89 MAINAI DOZESS 32, 250 119, 730 163, 846 75, 850 65, 245 75, 116 103, 624 109, 89 MAINAI DOZESS 37, 151 50, 304 36, 811 8, 507 18, 857 25, 978 44, 085 47, 00 MAINAI DOZESS 37, 141 37, 141 37, 141 37, 147 37, 1	AMAMAIODZÉSZ 7,74 9,361 3,697 10,386 2,003 4,100 3,302 9,849 AMAMAIODZÉSZ 36,715 23,391 15,641 22,341 10,671 20,167 32,429 30,795 AMAMAIODZÉSZ 36,715 23,393 15,641 22,341 10,671 20,167 32,429 30,795 AMAMAIODZÉSZ 18,342 10,645 5,272 3,3153 4,354 11,796 12,128 34,409 AMAMAIODZÉSZ 18,342 10,645 5,272 3,333 2,314 3,625 12,679 6,883 AMAMAIODZÉSZ 32,250 119,730 163,846 75,850 65,245 76,116 103,624 1096,697 AMAMAIODZÉSZ 32,250 119,730 163,846 75,850 65,245 76,116 103,624 1096,697 AMAMAIODZÉSZ 12,851 50,304 36,811 8,507 18,857 25,978 44,085 47,001 AMAMAIODZÉSZ 5,151 50,304 36,811 8,507 18,857 25,978 44,085 47,001 AMAMAIODZÉSZ 13,267 47,256 60,360,516 86,920 49,826 57,925 48,421 76,576 AMAMAIODZÉSZ 67,425 86,871 40,814 4,586 3,879 1,579 6,885 6,203 AMAMAIODZÉSZ 67,425 86,871 10,688 183,567 79,031 106,740 25,774 1,740 33,088 AMAMAIODZÉSZ 4,194 5,317 4,081 4,586 3,879 1,579 6,885 6,203 AMAMAIODZÉSZ 4,194 5,317 4,081 4,586 3,879 1,579 6,885 6,203 AMAMAIODZÉZZ 4,194 5,317 4,081 4,586 3,879 1,273 6,282 6,782 8 AMAMAIODZÉZZ 4,194 5,317 4,081 4,586 3,879 1,273 6,283 6,203 AMAMAIODZÉZZ 4,194 5,317 4,081 4,586 3,879 1,273 6,283 6,203 AMAMAIODZÉZZ 4,868 37 191,088 183,567 79,031 106,740 257,174 155,771 125,203 AMAMAIODZÉZZ 4,688 37,588 65,184 190,580 59,599 40,023 31,273 23,733 26,228 67,828 AMAMAIODZÉZZ 4,688 37,588 65,184 190,580 59,599 40,023 31,273 23,757 5,598 1 AMAMAIODZÉZZ 5,688 5,889 8,899 44,022 29,371 15,039 15,857 30,299 AMAMAIODZÉZZ 6,888 5,889 8,899 44,022 29,371 15,039 15,857 30,299 AMAMAIODZÉZZ 6,884 5,872 10,488 33,940 18,689 8,85,679 9,2471 84,01				21 795					
NAMBAIOOZESS	LAMAN 10027693 66.711 52.319 15.641 12.934 10.671 20.167 32.429 30.795 ALMAN 10027698 33.272 32.200 43.657 33.333 2.314 3.625 12.679 6.883 ALMAN 1002701 63.93 107.821 326.150 82.189 33.993 57.919 29.820 55.144 ALMAN 1002701 63.93 107.821 326.150 82.189 33.993 57.919 29.820 55.144 ALMAN 1002701 128.862 101.834 359.100 105.535 79.020 76.543 26.105 68.813 ALMAN 1002711 128.862 101.834 359.100 105.535 79.020 76.543 26.105 68.91 107.821 326.150 82.189 33.993 57.919 29.820 55.144 ALMAN 1002715 55.151 50.304 36.811 8.507 18.857 57.978 44.065 47.005									
NAMMA 1002779	AMAMAIODZ768 39 272 32 200 43 657 33 153 4 354 11 796 12 328 34 409									
LAMMATIODZ799 18.348 10.645 5.272 3.333 2.314 3.625 12.679 6.88 LAMMATIODZ701 66.193 107.821 326.150 82.189 33.991 57.919 29.820 55.141 LAMMATIODZ702 212.250 119.730 163.846 75.850 65.245 76.116 103.624 109.69 LAMMATIODZ712 55.151 50.304 38.9100 105.535 79.020 76.543 26.135 61.97 LAMMATIODZ712 55.151 50.304 38.811 8.507 18.857 25.978 44.085 47.00 LAMMATIODZ712 186.620 78.606 38.811 8.507 18.857 25.978 44.085 47.00 LAMMATIODZ712 128.620 78.606 38.01 8.50 21.356 39.333 49.740 33.08 LAMMATIODZ712 128.620 78.606 38.05.16 86.920 49.826 57.925 44.421 76.00 LAMMATIODZ712 128.620 78.606 380.516 86.920 49.826 57.925 44.421 76.00 LAMMATIODZ712 128.620 78.606 380.516 86.920 49.826 57.925 44.421 76.00 LAMMATIODZ712 4.194 5.317 4.081 4.586 3.859 39.1679 6.885 6.782 LAMMATIODZ722 4.55.508 63.239 134.784 49.359 17.238 32.733 26.228 67.22 LAMMATIODZ722 4.55.60 83.239 134.784 49.359 17.238 32.733 26.228 67.22 LAMMATIODZ723 45.508 83.239 134.784 49.359 17.238 32.733 26.228 67.22 LAMMATIODZ742 45.508 83.239 134.784 49.359 17.238 32.733 26.228 67.22 LAMMATIODZ742 45.508 83.239 134.784 49.359 17.238 32.733 26.228 67.22 LAMMATIODZ743 17.914 25.779 65.371 19.554 18.481 12.214 24.184 22.184 LAMMATIODZ744 17.914 68.871 19.088 183.567 79.031 108.740 257.774 156.771 125.28 LAMMATIODZ743 17.914 68.871 19.088 183.567 79.031 108.740 257.774 156.771 125.28 LAMMATIODZ745 4.60.93 65.485 18.49 40.02 2.93 79.116 3.957 9.800 1.039 7.99 LAMMATIODZ745 4.60.93 65.485 18.96 89.89 89.00 89.00 1.039 7.99 LAMMATIODZ745 4.60.93 65.485 18.99 40.02 2.93 79.116 3.957 9.800 1.039 7.99 LAMMATIODZ745 5.824 58.90 89.80 99.80 99.80 1.998 99.80 1.998 99.80 LAMMATIODZ745 65.245 88.90 99.755 5.507 5.640 9.500 11.988 99.70 LAMMATIODZ757 99.80	LIMINATIODZ789 18. 348 10. 645 5. 272 3. 333 2. 314 3. 625 12.679 6.883 UNINATIODZ780 28. 23. 250 119.7821 326.150 82. 189 33. 993 57. 919 39. 2820 55. 144 AARMATIODZ781 128. 862 101. 834 359. 100 105. 535 79. 920 75. 543 26. 135 61. 975 AARMATIODZ711 128. 862 101. 834 359. 100 105. 535 79. 920 75. 543 26. 135 61. 975 AARMATIODZ712 55. 151 50. 304 36. 811 8. 507 18. 857 25. 978 44. 0.85 47. 40. 854 AARMATIODZ716 32. 821 37. 741 37. 674 27. 554 13. 366 39. 383 49. 740 33. 088 AARMATIODZ716 18. 821 87. 741 37. 674 27. 554 13. 366 39. 383 49. 740 33. 088 AARMATIODZ727 18. 620 78. 806 360. 516 86. 920 49. 826 57. 925 48. 421 76. 78 AARMATIODZ727 18. 620 86. 85. 157 59. 116 53. 954 27. 853 31. 846 28. 039 37. 993 AARMATIODZ727 4. 194 5. 317 4. 081 4. 586 3. 879 11. 879 6. 885 6. 28. 039 AARMATIODZ728 455. 88 63. 239 134. 784 49. 369 17. 238 32. 733 26. 228 67. 828 AARMATIODZ728 45. 88. 71 191. 088 183. 567 79. 031 108. 740 257. 774 155. 771 126. 280 AARMATIODZ748 456. 871 191. 088 183. 567 79. 031 108. 740 257. 774 155. 771 126. 280 AARMATIODZ748 51. 3355 180. 986 171. 425 25. 271 3. 510 13. 742 11. 775 23. 747 AARMATIODZ748 45. 14. 967 8. 271 6. 239 9. 116 3. 957 9. 800 1. 039 7. 711 AARMATIODZ748 51. 3355 180. 986 171. 425 25. 271 3. 510 13. 742 11. 775 22. 747 AARMATIODZ758 26. 824 58. 122 104. 988 33. 940 18. 688 86. 579 9. 24. 71 18. 527 9. 11. 63. 98 91 AARMATIODZ758 27. 82. 335 7. 240 9. 756 5. 507 5. 640 9. 500 11. 958 9. 173 AARMATIODZ758 27. 82. 82. 82. 82. 82. 82. 82. 82. 82. 82	MAMMA 1002693	66.711	52. 339	15.641	32.934	10.671	20.167	32.429	30.795
LANDAL 1002769 18.348 10.645 5.272 3.333 2.314 3.625 12.679 5.88 LANDAL 1002708 212.250 119.730 163.846 75.850 65.245 76.116 103.624 109.69 MANENA 1002711 55.151 50.304 36.811 8.507 18.857 25.978 44.085 47.035 34.002712 55.151 50.304 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 44.085 47.033 36.811 8.507 18.857 25.978 25.978 44.085 47.033 36.811 8.507 18.858 25.979 25.818 47.033 36.818 36.978 38.03 49.740 33.08 36.818 36.978 38.03 36.978 36	LIMMATIODZ769 18. 348 10. 645 5. 272 3. 333 2. 314 3. 525 12. 679 6. 883 UMBATIODZ761 66. 193 107. 821 326.150 82. 193 33. 939 57. 919 9. 28. 820 584 134 AABATIODZ768 222. 253 119.730 163. 846 75. 850 65. 245 76. 116 103. 624 109. 697 AABATIODZ771 128. 862 101. 834 359. 100 105. 535 79. 020 75. 543 26. 135 61. 975 AABATIODZ771 128. 862 101. 834 359. 100 105. 535 79. 020 75. 543 26. 135 61. 975 AABATIODZ771 128. 862 1701. 834 359. 100 105. 535 79. 020 75. 543 26. 135 61. 975 AABATIODZ771 128. 862 77. 80. 036. 511 8. 507 18. 857 25. 978 44. 0.85 47. 031 0.84 100 0.85 12. 031 0.85 12. 0	JANNA1002698	39, 272	32, 200	43, 657	33, 153	4, 354	11.796	12, 328	34, 409
NAMAJOOZYOL 66,193 107,821 326,150 82,188 13,993 57,919 29,820 56,14 NAMAJOOZYOL 711 128,882 101,834 389,100 105,535 79,020 76,543 25,135 61,37 NAMAJOOZYOL 712 128,620 137,741 39,7674 23,554 13,666 39,383 49,740 33, NAMAJOOZYOL 32,825 80,000 300,516 86,920 49,826 57,925 48,421 76,57 NAMAJOOZYOL 128,620 78,060 360,516 86,920 49,826 57,925 48,421 76,57 NAMAJOOZYOL 128,620 78,060 360,516 86,920 49,826 57,925 48,421 76,57 NAMAJOOZYOL 128,630 36,317 4,081 4,585 3,879 1,679 6,885 6,20 NAMAJOOZYOL 48,508 63,239 134,784 49,359 17,238 32,733 26,628 61,22 NAMAJOOZYOL 48,508 63,239 134,784 49,359 17,238 32,733 26,6228 61,22 NAMAJOOZYOL 48,508 63,239 134,784 49,359 1,7238 32,733 26,6228 61,22 NAMAJOOZYOL 48,508 63,239 134,784 49,359 1,7238 32,733 26,6228 61,22 NAMAJOOZYOL 49,79 19,081 83,567 79,031 108,740 257,774 156,771 125,28 NAMAJOOZYOL 49,79 19,081 83,567 79,031 108,740 257,774 156,771 125,28 NAMAJOOZYOL 45 14,887 8,271 6,239 9,116 3,957 9,800 1,039 7,031 NAMAJOOZYOL 51,79 14 55,779 65,317 19,354 14,843 12,214 24,184 22,77 NAMAJOOZYOL 45 14,887 8,271 6,239 9,116 3,957 9,800 1,039 7,031 NAMAJOOZYOL 51,79 14 55,779 65,317 19,354 14,843 12,214 24,184 22,77 NAMAJOOZYOL 51,79 14 25,779 65,317 19,354 14,843 12,214 24,184 22,77 NAMAJOOZYOL 51,79 14 25,779 65,317 19,354 14,843 12,214 24,184 22,77 NAMAJOOZYOL 51,79 14 25,779 65,317 19,354 14,843 12,214 24,184 22,77 NAMAJOOZYOL 54 54,950 89,850 89,89 44,022 29,371 15,039 15,857 30,29 NAMAJOOZYOL 54 54,950 89,850 89,89 44,022 29,371 15,039 15,857 30,29 NAMAJOOZYOL 54 54,950 89,850 89,89 44,022 29,371 15,039 15,857 30,29 NAMAJOOZYOL 54 54,950 89,850 89,850 89,850 89,77 14,775 28,860 89,77 14,250 25,250 89,800 1,039 15,857 30,29 NAMAJOOZYOL 54 54,950 89,8	NAMALODOZYON 65, 193 107, 827 326, 150 82, 189 33, 993 57, 919 29, 820 55, 144								12 679	
NAMMA1002719 128.862 01.834 359.100 105.535 79.020 75.543 26.135 61.97 NAMMA1002711 128.862 01.834 359.100 105.535 79.020 75.543 26.135 61.97 NAMMA1002712 55.151 50.304 36.811 8.507 18.857 25.978 44.085 41.00 NAMMA1002713 128.621 37.741 37.674 23.554 13.366 39.333 49.740 33.08 NAMMA1002712 128.620 37.741 37.674 23.554 13.366 39.333 49.740 33.08 NAMMA1002721 128.620 78.060 360.516 86.920 49.825 57.925 48.421 76. NAMMA1002722 14.194 5.317 4.081 4.586 3.819 18.75 27.853 31.646 28.039 37.99 NAMMA1002727 4.194 5.317 4.081 4.586 3.819 1.679 6.885 6.782 NAMMA1002728 4.55.08 63.239 134.784 49.359 17.238 32.733 26.228 67.822 NAMMA1002742 466.871 191.088 183.567 79.031 108.740 257.374 1156.771 125.878 NAMMA1002743 17.911 25.779 65.317 19.354 14.843 12.214 24.184 22.27 NAMMA1002745 14.967 8.271 6.233 9.116 39.579 9.800 1.033 7.39 NAMMA1002746 14.967 8.271 6.233 9.116 39.579 9.800 1.033 7.39 NAMMA1002745 64.093 95.9489 171.425 25.271 9.150 39.579 9.800 1.033 7.017 9.354 14.843 12.214 24.184 22.27 NAMMA1002745 64.093 95.9489 19.956 59.599 40.023 33.273 23.675 53.99 NAMMA1002746 14.967 8.271 6.233 9.116 3.957 9.800 1.033 7.034 18.000 1.033 7.000 1.034	LAMMALODZYTOR 232, 250 119, 770 163, 846 75, 850 65, 245 76, 116 103, 824 109, 697 LAMMALODZYTOR 128, 862 101, 834 359, 100 105, 535 79, 9020 76, 543 25, 135 61, 975 LAMMALODZYTOR 232, 257 871 37, 741 37, 674 27, 554 13, 366 33, 383 49, 740 33, 088 LAMMALODZYTOR 128, 620 77, 68, 603 160, 516 86, 592 49, 825 57, 925 44, 085 47, 001 LAMMALODZYTOR 128, 620 77, 68, 603 160, 516 86, 592 49, 825 57, 925 44, 087 LAMMALODZYZOR 128, 620 77, 68, 603 160, 516 86, 592 49, 825 57, 925 44, 027 LAMMALODZYZOR 15, 64, 593 57, 59, 116 53, 954 27, 853 31, 646 28, 039 37, 993 LAMMALODZYZOR 41, 94 5, 317 4, 081 4, 586 3, 879 1, 679 6, 885 6, 202 LAMMALODZYZOR 45, 508 63, 239 134, 784 49, 359 17, 238 32, 733 26, 228 67, 822 LAMMALODZYZOR 45, 508 63, 239 134, 784 49, 359 17, 238 32, 733 26, 228 67, 822 LAMMALODZYZOR 45, 508 63, 239 134, 784 49, 359 17, 238 32, 733 26, 228 67, 822 LAMMALODZYZOR 47, 191 088 183, 557 79, 031 108, 740 257, 774 156, 777 126, 238 LAMMALODZYZOR 47, 191 088 183, 557 79, 031 108, 740 275, 774 156, 777 126, 238 LAMMALODZYZOR 47, 191 088 183, 557 79, 031 108, 740 275, 774 156, 777 126, 238 LAMMALODZYZOR 48, 53, 58, 68 67, 12, 239 9, 116 3, 957 9, 800 1, 038 7, 714 LAMMALODZYZOR 50, 50, 489 183, 849 44, 022 29, 371 15, 039 1, 10, 740 1, 10, 740 LAMMALODZYZOR 50, 50, 489 183, 499 44, 022 29, 371 15, 039 1, 10, 740 1,									
NAMARIO02712	ALMANIA1002715 128.862 101.814 359.106 105.515 79.020 76.543 25.135 61.975									
MANNAH 1002712	LAMMATODZ712 55. 151 50.304 36.811 8.507 18.857 25.978 44.085 47.001 ALMMATODZ718 32.821 37.741 37.674 23.554 13.366 39.383 49.740 33.088 AAMMATODZ721 128.520 78.060 360.516 86.920 49.826 57.925 48.421 76.576 AAMMATODZ727 4.194 5.317 4.081 4.586 3.879 1.679 5.885 31.646 28.039 37.933 AAMMATODZ727 4.194 5.317 4.081 4.586 3.879 1.679 6.885 6.203 AAMMATODZ728 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMATODZ728 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMATODZ728 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMATODZ728 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMATODZ724 486.871 191.088 183.567 79.031 108.740 257.734 156.771 16.728 AAMMATODZ725 4.060 70.772 65.1841 190.550 59.599 40.023 33.273 26.228 67.828 AAMMATODZ745 14.957 8.271 6.293 9.116 3.957 9.800 1.039 7.011 AAMMATODZ745 46.4093 69.489 189.499 44.022 29.77 AAMMATODZ745 46.4093 69.489 189.499 44.022 29.77 15.039 15.857 30.299 AAMMATODZ752 65.824 58.122 104.988 33.940 18.698 56.79 9.500 1.039 15.857 30.299 AAMMATODZ758 25.835 7.240 9.756 5.507 5.640 9.500 1.039 15.857 30.299 AAMMATODZ758 30.96 54.82 185.658 58.83 39.465 57.006 47.508 45.629 19.431 AAMMATODZ758 70.488 95.058 295.803 59.465 18.698 85.679 92.471 84.012 AAMMATODZ758 70.488 95.058 295.803 59.465 57.006 47.508 45.629 19.431 AAMMATODZ758 70.488 95.058 295.803 59.465 57.006 47.508 45.629 19.431 AAMMATODZ769 10.488 95.058 295.803 59.465 57.006 47.508 45.629 19.431 AAMMATODZ769 70.658 84.431 28.564 34.957 12.528 89.335 12.529 19.431 AAMMATODZ769 70.678 28.506 76.753 28.366 76.753 28.366 78.753 28.366 78.959 19.431 AAMMATODZ769 70.438 28.566 28.506 76.753 28.366 12.337 13.199 6.027 19.173 AAMMATODZ769 70.438 28.566 76.753 28.366 12.337 13.199 6.027 19.173 AAMMATODZ769 70.438 28.566 76.753 28.366 12.337 13.199 6.027 19.173 AAMMATODZ769 70.438 28.566 13.006 76.753 28.366 12.337 13.199 6.027 19.173 AAMMATODZ769 70.438 28.566 13.006 76.753 28.366 12.337 13.199 6.027 19.173 AAMMATODZ769 70.438 28.566 18.567 28.568	MAMMA1002708								
MANNA 1002776 32. 821 37. 741 37. 674 23. 554 13. 366 39. 383 49. 740 33. 084 144 144 144 145 145 145 145 145 145 14	LANDAL 1002716 32. 821 37. 741 37. 674 23. 554 13. 366 39. 383 49. 740 33. 088 43. MAINA 1002727 128. 620 78. 060 360. 516 86. 920 49. 825 57. 925 48. 421 76. 576 AABKAH 1002727 67. 425 45. 775 53. 116 53. 954 27. 853 31. 646 28. 039 37. 993 AABKAH 1002727 41. 94 5. 317 4. 081 4. 586 3. 879 1. 679 6. 885 6. 203 39 37. 993 AABKAH 1002728 45. 508 63. 239 134. 784 49. 359 17. 238 32. 733 26. 228 67. 828 AABKAH 1002728 45. 508 63. 239 134. 784 49. 359 17. 238 32. 733 26. 228 67. 828 AABKAH 1002742 48. 86. 871 191. 088 183. 567 79. 031 108. 740 257. 374 155. 771 125. 280 42. 40. 40. 40. 40. 40. 40. 40. 40. 40. 40	MAMMA1002711	128.862	101.834	359.100	105. 535		76. 543	<u>26.</u> 135	
LAIMAN 1002716 32. 821 37. 741 37. 674 23. 554 13. 366 39. 383 49. 740 33. 08. 44. MINIA 1002727 128. 620 78. 050 360. 516 86. 920 49. 826 57. 925 48. 821 76. 57 LAIMAN 1002727 178. 67. 425 45. 775 59. 116 53. 954 27. 853 31. 646 28. 039 37. 99 LAIMAN 1002728 45. 508 63. 239 314. 784 49. 359 17. 238 32. 733 32. 6228 67. 82 LAIMAN 1002728 45. 508 63. 239 314. 784 49. 359 17. 238 32. 733 32. 6228 67. 82 LAIMAN 1002727 44. 945 52. 86. 86. 871 191. 088 183. 567 79. 031 108. 740 257. 774 156. 771 125. 28 LAIMAN 1002742 46. 871 191. 088 183. 567 79. 031 108. 740 257. 774 156. 771 125. 28 LAIMAN 1002743 17. 914 25. 779 65. 317 19. 354 14. 843 12. 214 24. 184 22. 27 LAIMAN 1002744 70. 172 65. 184, 190. 550 59. 599 40. 023 33. 273 23. 675 53. 99 LAIMAN 1002744 53. 355 180. 986 171. 425 25. 277 3. 510 13. 742 11. 775 23. 74 LAIMAN 1002748 53. 355 180. 986 171. 425 25. 277 3. 510 13. 742 11. 775 23. 74 LAIMAN 1002758 25. 835 7. 240 9. 756 5. 507 5. 640 9. 500 11. 968 9. 17 LAIMAN 1002758 25. 835 7. 240 9. 756 5. 507 5. 640 9. 500 11. 968 9. 17 LAIMAN 1002765 10. 88 9. 568 95. 803 59. 645 52. 006 47. 508 45. 529 11. 56. 98 LAIMAN 1002775 9. 652 248. 038 1.22 104. 988 33. 940 18. 698 86. 679 9. 2471 84. 01 LAIMAN 1002765 81. 926 54. 425 185. 685 56. 538 25. 534 30. 254 22. 519 36. 21 LAIMAN 1002775 9. 652 248. 038 1. 136 106. 297 36. 324 95. 235 52. 022 923. 91 LAIMAN 1002775 9. 652 248. 038 1. 136 106. 297 36. 324 95. 235 52. 022 923. 91 LAIMAN 1002775 9. 652 248. 038 1. 136 106. 297 36. 324 95. 235 52. 022 923. 91 LAIMAN 1002775 9. 652 248. 038 1. 136 106. 297 36. 324 95. 235 52. 022 923. 91 LAIMAN 1002785 10. 41. 41. 41. 41. 41. 41. 41. 41. 41. 41	LANIMA 1002716 32. 821 37.741 37.674 23.554 13.366 19.383 49.740 33.088 AIMMA 1002727 128.620 78.060 360.516 85.920 49.826 57.925 48.421 76.576 AAMMA 1002727 67.425 45.775 59.116 53.954 27.853 31.648 28.039 37.993 AAMMA 1002727 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMA 1002727 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMA 1002728 45.508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AAMMA 1002742 48.68.871 191.088 183.567 79.031 108.740 257.374 155.771 125.236 AAMMA 1002743 17.914 25.779 65.317 19.354 14.84 31 12.214 24.184 22.277 AAMMA 1002743 17.914 25.779 65.317 19.354 14.84 31 12.214 24.184 22.277 AAMMA 1002744 70.172 65.184, 190.550 59.599 40.023 33.273 23.675 33.991 AAMMA 1002745 45.355 88.271 6.293 9.116 3.957 9.800 1.039 7.011 AAMMA 1002745 45.355 88.291 6.293 9.116 3.957 9.800 1.039 7.011 AAMMA 1002745 45.355 89.595 171.425 25.271 3.510 13.742 11.775 23.747 AAMMA 1002754 64.093 69.489 189.499 44.022 29.371 15.039 15.857 30.259 14.444 10.2146 14.04 14	MAHMA1002712	55, 151	50, 304	36.811	8.507	18.857	25.978	44.085	47.001
WANNAH 1002721 128, 520 78, 050 360, 516 36, 320 49, 826 57, 925 48, 421 76, 57	AMMATOOZ721 128.520 78.060 360.516 36. 920 49.825 57.925 48.421 76.576 AMMATOOZ727 7.4794 55.775 59.116 53.954 27.853 31.646 28.039 37.993 AMMATOOZ727 4.194 5.317 4.081 4.586 3.879 1.679 6.885 6.203 AMMATOOZ728 4.5508 63.239 134.784 49.359 17.238 32.733 26.228 67.828 AMMATOOZ728 4.85.871 191.088 183.567 79.031 108.740 257.374 156.771 126.280 AMMATOOZ724 486.871 191.088 183.567 79.031 108.740 257.374 156.771 126.280 AMMATOOZ744 70.172 65.184, 190.550 59.599 40.023 33.273 22.675 53.991 AMMATOOZ745 14.857 8.271 6.293 9.116 3.957 9.800 1.038 7.011 AMMATOOZ746 14.857 8.271 6.293 9.116 3.957 9.800 1.038 7.011 AMMATOOZ745 64.093 69.489 189.499 44.022 92.971 15.039 15.857 30.299 AMMATOOZ756 64.093 69.489 189.499 44.022 92.971 15.039 15.857 30.299 AMMATOOZ756 25.835 7.240 9.756 5.507 5.640 9.500 11.968 9.173 AMMATOOZ756 26.824 88.122 104.988 13.940 14.698 83.940 94.923 10.254 12.44 4.019.868 9.173 AMMATOOZ757 50 50.978 9.602 33.997 9.800 11.968 9.173 AMMATOOZ757 50 9.062 33.997 9.878 5.366 12.293 19.431 57.791 AMMATOOZ767 50 9.062 33.997 9.878 5.366 12.293 19.431 57.791 AMMATOOZ775 51.236 54.425 185.685 36.838 25.634 30.254 22.519 36.212 AMMATOOZ775 51.236 54.425 185.685 36.838 25.634 30.254 22.519 36.212 AMMATOOZ775 51.236 37.084 125.540 30.088 37.975 22.272 39.91.11 AMMATOOZ775 51.236 37.084 125.540 30.088 37.975 22.275 30.292 39.91 AMMATOOZ775 51.236 37.084 125.540 30.088 37.975 22.242 25.695 33.531 AMMATOOZ780 23.190 24.577 75.773 29.564 12.337 13.199 6.027 19.174 AMMATOOZ780 75.75 12.266 75.828 30.086 77.788 18.969 30.126 13.855 9.331 47.78 29.564 12.337 13.199 6.027 19.174 AMMATOOZ780 75.75 12.266 75.75 75.75 28.266 12.337 12.258 19.351 10.186 19.521 AMMATOOZ780 75.75 12.266 75.75 75.75 12.75 12.258 19.355 12.259 12.259 12.359 12.259 12.359 12.259 12.359 12.259 12.359 12.259 12.359 12.259 12.359 12.						13, 366	39, 383	49.740	33.088
MAJMAN 10027723	ALMMA 1002723 67, 425 45, 775 59, 116 53, 954 27, 853 31, 646 28, 0.39 37, 933 ALMMA 1002727 45, 508 63, 239 134, 184 49, 389 17, 238 32, 733 26, 228 67, 828 ALMMA 1002742 48, 68, 871 191, 088 183, 567 79, 031 108, 740 257, 374 155, 771 126, 236 ALMMA 1002743 17, 914 25, 779 65, 817 79, 931 108, 740 257, 374 155, 771 126, 236 ALMMA 1002744 70, 172 65, 184, 190, 550 59, 599 40, 0.23 32, 273 23, 675 33, 991 ALMMA 1002745 14, 967 8, 271 6, 293 9, 116 3, 957 9, 800 1, 0.39 7, 0.11 ALMMA 1002745 53, 355 180, 986 171, 425 25, 271 3, 510 13, 742 11, 775 23, 747 ALMMA 1002745 46, 0.93 69, 489 189, 499 44, 022 29, 371 15, 0.39 15, 857 30, 299 ALMMA 1002755 82, 58, 35 7, 240 9, 756 5, 507 5, 640 9, 500 11, 968 9, 340 ALMMA 1002765 81, 326 54, 425 185, 685 36, 838 25, 634 30, 254 22, 517 ALMMA 1002765 81, 326 54, 425 185, 685 36, 838 27, 634 30, 254 22, 519 ALMMA 1002777 92, 652 248, 038 91, 136 106, 297 33, 539 15, 867 29, 91 ALMMA 1002778 92, 652 248, 038 91, 136 106, 297 34, 324 25, 529 31, 53, 540 ALMMA 1002779 92, 652 248, 038 91, 136 106, 297 36, 324 35, 252 35, 92, 92, 91 ALMMA 1002778 75, 12, 36 37, 084 125, 540 30, 088 37, 975 21, 242 25, 595 22, 939 14 ALMMA 1002780 27, 192, 652 248, 038 91, 136 106, 297 36, 324 35, 235 32, 022 99, 91 ALMMA 1002780 27, 192, 652 28, 506 67, 753 28, 366 9, 933 47, 769 23, 31, 31, 340 ALMMA 1002780 27, 192, 652 28, 506 67, 753 28, 366 9, 933 47, 769 23, 31, 31, 31, 34, 34, 34, 34, 34, 34, 34, 34, 34, 34									
NAMMA1002727	LAMMA10027727									
MAMMA1002718	LAMMA1002743									
Mamma1002742 486.871 91.088 183.567 79.031 108.740 257.374 156.771 125.28	LAMMA1002742									
MAMMA1002743	HAMMA1002743	MAMMA1002728	45. 508	63. 239	134.784	49.369	17. 238		26. 228	
NAMMA1002743	MANIMA 1002743	MAMMA1002742	486, 871	191.088	183.567	79.031	108.740	257. 374	156.771	126.280
MAMMA1002744	MANIMA 1002744					19.354	14.843	12.214	24. 184	22, 277
MANIMA 1002746	LAMMAT 1002746 14.967 8.271 6.293 9.116 3.957 9.800 1.039 7.011 LAMMAT 1002748 53.355 180.966 171.425 25.271 3.510 13.742 11.775 23.747 LAMMAT 1002754 64.093 69.489 188.499 44.022 29.371 15.039 15.857 30.299 LAMMAT 1002752 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 LAMMAT 1002752 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 LAMMAT 1002752 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 LAMMAT 1002752 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 LAMMAT 1002752 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 LAMMAT 1002752 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 LAMMAT 1002752 65.825 248.038 91.36 105.297 36.324 95.235 52.022 929.910 LAMMAT 10027771 92.652 248.038 91.36 105.297 36.324 95.235 52.022 929.910 LAMMAT 10027771 92.652 248.038 91.36 105.297 36.324 95.235 52.022 929.910 LAMMAT 10027780 23.190 24.572 73.778 29.564 12.337 31.199 6.027 19.178 LAMMAT 10027780 23.190 24.572 73.778 29.564 12.337 31.199 6.027 19.178 LAMMAT 1002780 23.190 24.572 73.778 29.564 12.337 31.199 6.027 19.178 LAMMAT 1002780 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.835 LAMMAT 1002780 88.4 31 28.564 34.957 32.528 49.335 29.125 31.088 19.921 LAMMAT 1002805 88.4 31 28.564 34.957 32.528 49.335 29.125 31.050 33.944 LAMMAT 1002807 64.374 42.471 124.050 39.454 51.288 34.538 23.265 46.125 LAMMAT 1002817 8.719 10.443 6.527 4.336 14.707 19.459 22.550 33.531 LAMMAT 1002817 8.719 10.43 6.527 4.336 14.707 19.459 22.550 33.531 LAMMAT 1002817 8.719 10.43 6.527 4.336 14.707 19.459 22.550 33.531 LAMMAT 1002833 90.875 71.138 23.7238 50.346 44.689 47.222 25.094 46.036 LAMMAT 1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.22 LAMMAT 1002833 90.875 71.33 13.566 35.466 14.707 19.459 22.550 33.531 LAMMAT 1002835 84.568 22.248 88.823.244 28.102 14.935 9.604 12.597 16.302 12.704 LAMMAT 1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.22 LAMMAT 1002843 98.706 53.519 151.675 23.902 33.008 77.477 54.600 33.856 33.856 33.590 18.646									
MANIATIO02748	MANIMA 1002748									
MANIMA 1002754	NAMIA 1002754 64. 093 69. 489 189. 499 44. 022 29. 371 15. 039 15. 857 30. 299 NAMIA 1002758 25. 835 7. 240 9.756 5. 507 5. 640 9. 500 11. 968 9. 173 NAMIA 1002762 65. 824 58. 122 104. 988 33. 940 18. 698 86. 679 92. 471 84. 012 NAMIA 1002764 104. 828 95. 058 295. 803 59. 465 52. 006 47. 508 45. 629 48. 337 NAMIA 1002765 81. 926 54. 425 185. 685 56. 838 25. 634 30. 254 22. 519 36. 212 NAMIA 1002769 20. 078 9. 052 33. 997 9. 878 15. 366 12. 293 19. 431 15. 797 NAMIA 1002777 51. 236 37. 084 125. 540 30. 088 37. 975 21. 242 25. 693 40. NAMIA 1002778 52. 636 37. 084 125. 540 30. 088 37. 975 21. 242 25. 693 24. 95. 235 12. 022 929. 910 NAMIA 1002778 51. 236 37. 084 125. 540 30. 088 37. 975 21. 242 25. 693 24. 95. 235 12. 022 929. 910 NAMIA 1002778 52. 636 37. 084 125. 540 30. 088 37. 975 21. 242 25. 693 24. 95. 235 12. 022 929. 910 NAMIA 1002780 23. 190 24. 572 73. 778 29. 564 12. 337 13. 199 6. 027 19. 175 NAMIA 1002780 25. 198 18. 430 30. 126 38. 38. 94 12. 340 12. 34			8. 271						
MAMMA1002765	NAMMATOOZ765 25.835 7.240 9.756 5.507 5.640 9.500 11.968 9.173 NAMMATOOZ762 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.012 NAMMATOOZ765 81.926 54.425 185.685 55.803 59.465 52.006 47.508 45.629 48.337 NAMMATOOZ765 81.926 54.425 185.685 56.838 25.634 30.254 22.519 36.212 NAMMATOOZ765 81.926 54.425 185.685 56.838 25.634 30.254 22.519 36.212 NAMMATOOZ767 92.0078 9.062 33.997 9.878 15.366 12.293 19.431 15.79 NAMMATOOZ777 92.652 248.038 91.136 105.297 36.324 95.235 52.022 929.910 NAMMATOOZ778 51.236 37.084 125.540 30.088 37.975 21.242 25.695 24.381 NAMMATOOZ780 23.190 24.572 73.778 29.564 12.337 13.199 6.027 19.175 NAMMATOOZ782 76.728 28.066 76.753 28.366 26.053 26.045 13.885 33.944 NAMMATOOZ785 17.412 3.178 14.907 9.264 2.359 6.615 10.186 19.921 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.494 19.700 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 NAMMATOOZ805 84.431 28.564 34.957 25.258 49.335 29.125 31.705 30.488 NAMMATOOZ805 76.4374 42.471 124.060 39.454 51.288 34.538 23.255 46.125 NAMMATOOZ805 77.604 20.918 18.590 24.282 25.90 33.531 NAMMATOOZ805 84.431 31.853 13.9150 28.8560 58.851 17.5492 49.491 111.835 311.632 133.131 NAMMATOOZ805 85.519 15.675 23.002 28.250 07.782 25.904 46.081 NAMMATOOZ805 97.634 37.795 15.675 23.002 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.802 37.797 18.804 37.797 18.802 37.797 18.804 37.797 18.904 37.797 18.804 37.797 18.804 37.797 18.804 37.797 18.804 37			180.966						
MAMMA1002762 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.01	MAMMATIO02762 65. 824 58. 122 104. 988 33. 940 18. 698 86.679 92. 471 84. 012 MAMMATIO02765 104. 828 95. 058 295. 803 59. 465 52. 006 47. 508 45. 629 48. 337 MAMMATIO02765 81. 926 54. 425 185. 685 56. 838 25. 634 30. 254 42. 519 36. 212 MAMMATIO02769 20. 078 9. 062 33. 997 9. 878 15. 366 12. 293 19. 431 15. 799 MAMMATIO027775 51. 235 37. 084 125. 540 30. 088 37. 975 21. 242 25. 695 24. 387 MAMMATIO027775 51. 236 37. 084 125. 540 30. 088 37. 975 21. 242 25. 695 24. 387 MAMMATIO02778 02. 31. 90 24. 572 73. 778 29. 564 12. 337 13. 199 6. 027 19. 175 MAMMATIO02779 76. 728 28. 066 76. 753 28. 366 26. 053 26. 045 13. 885 33. 944 MAMMATIO02795 77. 142 3. 178 14. 907 9. 264 2. 359 6. 615 10. 186 19. 921 MAMMATIO02796 28. 596 28. 390 48. 340 13. 930 16. 360 14. 274 13. 494 19. 700 MAMMATIO02806 84. 431 28. 564 34. 957 22. 528 49. 335 29. 125 31. 705 30. 485 MAMMATIO02807 64. 374 42. 471 124. 060 39. 454 51. 288 34. 538 23. 265 46. 12. MAMMATIO02807 76. 719 10. 443 6. 527 4. 036 11. 155 2. 240 8. 038 51. 11 MAMMATIO02807 77 87 99. 133. 566 28. 366 14. 707 19. 459 22. 590 33. 531 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 12 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 12 MAMMATIO02808 78. 488 23. 244 28. 102 14. 935 9. 604 12. 597 16. 302 12. 70 MAMMATIO02808 79. 71. 718 237. 238 50. 346 14. 689 47. 222 25. 094 46. 084 MAMMATIO02808 79. 71. 718 237. 238 50. 346 14. 689 47. 222 25. 094 46. 084 MAMMATIO02808 79. 71. 718 237. 238 50. 346 14. 689 79. 728 15. 728 15. 728 18. 728 19.									
MAMMA1002762 65.824 58.122 104.988 33.940 18.698 86.679 92.471 84.01	MAMMATIO02762 65. 824 58. 122 104. 988 33. 940 18. 698 86.679 92. 471 84. 012 MAMMATIO02765 104. 828 95. 058 295. 803 59. 465 52. 006 47. 508 45. 629 48. 337 MAMMATIO02765 81. 926 54. 425 185. 685 56. 838 25. 634 30. 254 42. 519 36. 212 MAMMATIO02769 20. 078 9. 062 33. 997 9. 878 15. 366 12. 293 19. 431 15. 799 MAMMATIO027775 51. 235 37. 084 125. 540 30. 088 37. 975 21. 242 25. 695 24. 387 MAMMATIO027775 51. 236 37. 084 125. 540 30. 088 37. 975 21. 242 25. 695 24. 387 MAMMATIO02778 02. 31. 90 24. 572 73. 778 29. 564 12. 337 13. 199 6. 027 19. 175 MAMMATIO02779 76. 728 28. 066 76. 753 28. 366 26. 053 26. 045 13. 885 33. 944 MAMMATIO02795 77. 142 3. 178 14. 907 9. 264 2. 359 6. 615 10. 186 19. 921 MAMMATIO02796 28. 596 28. 390 48. 340 13. 930 16. 360 14. 274 13. 494 19. 700 MAMMATIO02806 84. 431 28. 564 34. 957 22. 528 49. 335 29. 125 31. 705 30. 485 MAMMATIO02807 64. 374 42. 471 124. 060 39. 454 51. 288 34. 538 23. 265 46. 12. MAMMATIO02807 76. 719 10. 443 6. 527 4. 036 11. 155 2. 240 8. 038 51. 11 MAMMATIO02807 77 87 99. 133. 566 28. 366 14. 707 19. 459 22. 590 33. 531 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 11 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 12 MAMMATIO02807 78. 719 10. 443 6. 527 4. 036 1. 155 2. 240 8. 038 51. 12 MAMMATIO02808 78. 488 23. 244 28. 102 14. 935 9. 604 12. 597 16. 302 12. 70 MAMMATIO02808 79. 71. 718 237. 238 50. 346 14. 689 47. 222 25. 094 46. 084 MAMMATIO02808 79. 71. 718 237. 238 50. 346 14. 689 47. 222 25. 094 46. 084 MAMMATIO02808 79. 71. 718 237. 238 50. 346 14. 689 79. 728 15. 728 15. 728 18. 728 19.	MAMMA1002758	25, 835	7, 240	9.756	5. 507	5. 640	9, 500	11.968	9. 173
MAIMAT1002764	MAJMAN 1002786 104.828 95.058 295.803 59.465 52.006 47.508 45.629 48.337				104.988	33.940	18, 698		92, 471	84.012
MAIMAT1002765	MAIMATIOD2765									
MAIMAT1002769	MANIMATO02765 20.078 9.062 33.997 9.878 15.366 12.293 19.431 15.797									
MAIMA1002771 92.652 248.038 91.136 106.297 36.324 95.235 52.022 929.91	MAMMA1002771 92.652 248.038 91.136 106.297 36.324 95.235 52.022 929.910									
MAMMA1002785 51.236 37.084 125.540 30.088 37.975 21.242 25.695 24.38 MAMMA1002780 23.190 24.572 73.778 29.564 12.337 13.199 6.027 19.78 MAMMA1002782 67.728 28.066 76.753 28.366 26.053 26.045 13.885 33.94 MAMMA1002795 17.412 3.178 14.907 9.264 2.359 6.615 10.186 19.92 MAMMA1002795 28.596 28.390 48.340 13.930 16.360 14.274 13.494 19.70 MAMMA10027805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.43 MAMMA10027806 84.431 28.564 34.957 32.528 49.335 29.125 31.705 30.48 MAMMA10027807 64.374 42.471 124.080 39.454 51.288 34.538 23.265 46.12 MAMMA10027807 64.374 42.471 124.080 39.454 51.288 34.538 23.265 46.12 MAMMA10027807 64.374 42.471 124.080 39.454 51.288 34.538 23.265 46.12 MAMMA10027807 64.374 42.471 124.080 39.454 51.288 34.538 23.265 46.12 MAMMA10027807 64.374 42.471 124.080 39.454 51.288 34.538 23.265 46.12 MAMMA10027807 64.374 42.471 14.605 7.416 9.432 15.038 51.1 MAMMA10027807 64.374 42.471 42.050 39.454 51.288 34.538 23.265 46.12 43.481	MAMMA1002775 51.236 37.084 125.540 30.088 37.975 21.242 25.695 24.387 MAMMA1002780 23.190 24.572 73.778 29.564 12.337 13.199 6.027 19.175 MAMMA1002780 76.728 28.066 76.753 28.366 26.053 26.045 13.885 33.944 MAMMA1002780 17.412 3.178 14.907 9.264 2.359 6.615 10.186 19.921 MAMMA1002795 17.412 3.178 14.907 9.264 2.359 6.615 10.186 19.921 MAMMA1002805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.432 MAMMA1002806 84.431 28.564 34.957 32.528 49.335 29.125 31.705 30.488 MAMMA1002807 64.374 42.471 124.060 39.454 51.288 34.538 23.265 46.125 MAMMA1002807 64.374 42.471 124.060 39.454 51.288 34.538 23.265 46.125 MAMMA1002807 64.374 42.471 124.060 39.454 51.288 34.538 23.265 46.125 MAMMA1002817 8.719 10.443 6.527 4.036 11.155 2.240 8.038 11.125 MAMMA1002807 64.374 42.471 14.605 7.416 9.432 16.038 5.11 MAMMA1002807 64.374 42.471 14.4060 39.454 51.288 34.538 23.265 46.125 MAMMA1002817 8.719 10.443 6.527 4.036 11.155 2.240 8.038 11.125 MAMMA1002830 91.438 212.662 185.761 75.492 49.491 111.835 311.632 133.133 MAMMA1002830 99.875 71.138 237.238 50.346 44.669 47.222 25.590 33.13.133 MAMMA1002833 90.875 71.138 237.238 50.346 44.669 47.222 25.994 46.086 MAMMA1002838 84.752 56.692 166.200 49.694 30.237 32.930 11.628 26.416 MAMMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.16 MAMMA1002857 77.604 209.913 235.780 167.479 18.190 24.282 20.456 19.401 13.72 MAMMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.091 MAMMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 74.77 5.467 3.406 MAMMA1002880 65.375 102.643 253.035 92.062 91.774 46.567 74.77 5.467 3.294 MAMMA1002887 77.604 209.913 235.780 167.189 50.200 77.282 50.944 50.081 MAMMA1002888 13.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.091 MAMMA1002887 77.604 209.913 235.780 167.497 18.190 24.282 201.758 152.946 94.161 MAMMA1002887 87.033 19.03 38.206 52.831 26.325 19.774 55.667 34.650 30.655 19.401 33.72 MAMMA1002887 88.545 53.572 210.646 36.086 31.508 88.271 33.675 5.11 MAMMA1002888 65.375 102.643 38.977 25.333 27.378 18.964									
MAMMA1002780 23.190 24.572 73.778 29.564 12.337 13.199 6.027 19.77	MAMMA1002780 23.190 24.572 73.778 29.564 12.337 13.199 6.027 19.175 MAMMA1002782 76.728 28.066 76.753 28.366 26.053 26.045 13.885 33.944 MAMMA1002796 28.596 28.390 48.340 13.930 16.360 14.274 13.494 19.705 MAMMA1002805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.433 MAMMA1002806 84.431 28.564 34.957 32.528 49.335 29.125 31.705 30.485 MAMMA1002807 64.374 42.471 124.060 39.455 1288 34.538 23.265 46.125 MAMMA1002807 64.374 42.471 124.060 39.454 51.288 34.538 23.265 46.125 MAMMA1002814 28.078 31.573 133.666 36.466 14.707 19.459 22.590 33.533 MAMMA1002817 8.719 10.443 6.527 4.036 1.155 2.240 8.038 11.121 MAMMA1002800 15.173 5.049 24.747 14.605 7.416 9.432 16.038 5.11 MAMMA1002830 91.438 212.662 185.761 75.492 49.491 111.835 311.632 133.137 MAMMA1002833 90.875 71.138 237.238 50.346 44.689 47.222 25.094 46.080 MAMMA1002838 84.752 56.692 166.200 49.694 30.237 32.930 11.628 26.41 MAMMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.161 MAMMA1002857 77.604 209.913 235.780 167.148 50.207 178.228 129.737 278.80 MAMMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.091 MAMMA1002858 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002858 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MAMMA1002858 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.851 MAMMA1002859 85.453 22.923 80.058 19.164 22.993 25.241 45.735 80.077 23.10 MAMMA1002859 85.4	MAMMA1002771								
MAMMA1002780	MAMMA1002780 23.190 24.572 73.778 29.564 12.337 13.199 6.027 19.175 MAMMA1002785 76.728 28.066 76.753 28.366 26.053 26.045 13.885 33.944 MAMMA1002795 17.412 3.178 14.907 9.264 2.359 6.615 10.186 19.921 MAMMA1002796 28.596 28.390 48.340 13.930 16.360 14.274 13.494 19.705 MAMMA1002805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.432 MAMMA1002806 84.431 28.564 34.957 32.528 49.335 29.125 31.705 30.485 MAMMA1002807 64.374 42.471 124.060 39.455 1.288 34.538 23.265 46.125 MAMMA1002814 28.078 31.573 133.666 36.466 14.707 19.459 22.590 33.533 MAMMA1002817 8.719 10.443 6.527 4.036 11.55 2.240 8.038 11.121 MAMMA1002807 15.173 5.049 24.747 14.605 7.416 9.432 16.038 5.11 MAMMA1002830 91.438 212.662 185.761 75.492 49.491 111.835 311.632 133.135 MAMMA1002830 90.875 71.138 237.238 50.346 44.689 47.222 25.094 46.086 MAMMA1002832 84.752 56.692 166.200 49.694 30.237 32.930 11.622 133.135 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MAMMA1002844 311.853 319.050 65.856 66.881 72.282 201.758 152.946 MAMMA1002845 4.464 5.631 16.258 13.028 3.035 26.112 45.735 86.883 51.98 MAMMA1002845 5.56.692 166.200 49.694 30.237 32.3930 11.628 26.411 MAMMA1002845 5.56.692 166.200 49.694 30.237 32.3930 11.628 26.411 MAMMA1002847 13.899 319.730 662.654 523.500 84.144 532.413 382.518 1000.091 MAMMA1002885 85.375 102.643 253.035 92.062 91.774 46.567 38.439 58.466 MAMMA1002885 85.375 102.643 253.035 92.062 91.774 46.567 38.439 58.466 MAMMA1002887 20.954 16.542 18.160 22.628 23.110	MAMMA1002775	51.236	37. 084	125. 540	30.088	37. 975	21. 242	25, 695	24. 387
MANIMATO02782	MANIMAT1002782	MANMA1002780	23, 190	24.572		29.564	12, 337	13, 199	6.027	19. 175
MANIMATOD2895 17. 412 3. 178 14. 907 9. 264 2. 359 6. 615 10. 186 19. 92	MANMAIO02805									
MANMAIO02805 28. 596 28. 390 48. 340 13. 930 16. 360 14. 274 13. 494 19. 70	MARMA1002805 25.198 16.430 30.126 13.856 9.933 47.769 23.312 13.434 13.494 19.705									
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NAMMA1002817 8.719 10.443 6.527 4.036 1.155 2.240 8.038 11.12	MANNA1002817 8.719 10.443 6.527 4.036 1.155 2.240 8.038 11.12	MAMMA1002814	28.078	31, 573	133,666	36.456	14.707	19,459	22.590	33. 539
NAMMA1002830 15.173 5.049 24.747 14.605 7.416 9.432 16.038 5.11 NAMMA1002830 91.438 212.662 185.761 75.492 49.491 111.835 311.632 133.13 NAMMA1002833 90.875 71.138 237.238 50.346 44.689 47.222 25.094 46.08 NAMMA1002835 28.488 23.244 28.102 14.935 9.604 12.597 16.302 12.70 NAMMA1002838 84.752 56.692 166.200 49.694 30.237 32.930 11.628 26.41 NAMMA1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.22 NAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.455 19.401 13.77 NAMMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.16 NAMMA1002845 4.464 5.631 16.258 13.028 3.642 8.306 5.338 22.84 NAMMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.86 NAMMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.05 NAMMA1002868 85.375 102.643 253.035 92.062 91.774 46.567 38.439 58.44 NAMMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.81 NAMMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 NAMMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 NAMMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 NAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 NAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 NAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 NAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 NAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 NAMMA1002889 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 NAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 38.649 41.0 NAMMA1002889 58.445 53.672 210.646	MANNA1002830 15.173 5.049 24.747 14.605 7.416 9.432 16.038 5.11								8.038	11, 128
MANMA1002830 91.438 212.662 185.761 75.492 49.491 111.835 311.632 133.13 MANMA1002833 90.875 71.138 237.238 50.346 44.689 47.222 25.094 46.08 MANMA1002835 28.488 23.244 28.102 14.935 9.604 12.597 16.302 12.70 MANMA1002838 84.752 56.692 166.200 49.694 30.237 32.930 11.628 26.41 MANMA1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.22 MANMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.77 MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.16 MANMA1002845 4.464 5.631 16.258 13.028 3.642 8.306 5.338 22.84 MANMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.86 MANMA1002853 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.09 MANMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.91 MANMA1002868 85.375 102.643 253.035 92.062 91.774 46.567 38.439 58.44 MANMA1002875 20.954 16.542 18.160 26.628 23.110 21.099 24.952 32.94 MANMA1002875 20.954 16.542 18.160 26.628 23.110 21.099 24.952 32.94 MANMA1002875 20.954 16.542 18.160 26.628 23.110 21.099 24.952 32.94 MANMA1002875 20.954 16.542 18.160 26.628 23.110 21.099 24.952 32.94 MANMA1002875 37.255 55.154 238.977 25.333 27.378 18.964 34.053 52.4 MANMA1002887 33.550 7.099 12.119 12.813 15.447 20.107 22.3 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 33.649 41.0 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 33.649 41.0 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 33.649 41.0 MANMA1002887 53.455 53.672 210.646 36.086 31.508 31.508 36.186 13.729 35.7 MANMA1002893 58.445 53.672 210.646 36.086 31.508 31.508 36.186 13.729 35.7 MANMA1002893 76.469 18.593 25.600 5.864 9.	MANNA1002830 91.438 212.662 185.761 75.492 49.491 111.835 311.632 133.13; MANNA1002833 90.875 71.138 237.238 50.346 44.689 47.222 25.094 46.081 MANNA1002835 28.488 23.244 28.102 14.935 9.604 12.597 16.302 12.709 MANNA1002838 84.752 56.692 166.200 49.694 30.237 32.930 11.628 26.410 MANNA1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.222 MANNA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MANNA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.160 MANNA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MANNA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.09 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA100287 30.056 30.85 30.058									
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HAMMA1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.22 HAMMA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.77 HAMMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.16 HAMMA1002845 4.464 5.631 16.258 13.028 3.642 8.306 5.338 22.84 HAMMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.86 HAMMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.05 HAMMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.91 HAMMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 38.439 58.41 HAMMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.81 HAMMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 HAMMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 HAMMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 HAMMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 HAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 HAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 HAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 38.649 41.0 HAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 38.649 41.0 HAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 38.649 41.0 HAMMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 38.649 41.0 HAMMA1002889 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 HAMMA1002883 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2 HAMMA1002883 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2 HAMMA1002883 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2 HAMMA1002893 76.469 18.593 25.600 5.864	MANNA1002842 98.706 53.519 151.675 23.902 32.033 41.236 27.950 47.22 MANNA1002843 76.343 31.051 107.479 18.190 24.282 30.456 19.401 13.72 MANNA1002845 4.464 5.631 16.258 13.028 3.642 8.306 5.338 22.84 MANNA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MANNA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.09 MANNA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.98 MANNA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.98 MANNA1002868 85.375 102.643 253.035 92.062 91.774 46.567 38.439 58.46 MANNA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANNA1002879 23.352 14.773 9.446 6.359 8.506 13.275 30.077 23.10 MANNA1002879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.10 MANNA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 MANNA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.41 MANNA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.546 36.086 31.508 36.186 13.729 35.74 MANNA1002887 45.505 7.809 7.546 36.086 31.508 36.186 13.729 35.74 MANNA1002887 45.505 7.809 7.546 36.086 31.508 36.186 13.729 35.74 MANNA1002887 45.505 7.809 7.546 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.546 7.024 9.968 8.271 13.675 5.11 MANNA1002887 45.505 7.809 7.546 7.024 9.968 8.									
MAIMA1002843 76.343 31.051 107.479 18.190 24.282 30.455 19.401 13.72	MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.161 MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.161 MANMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MANMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.091 MANMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.98 MANMA1002868 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002875 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.10 MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 MANMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.41 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.546 6.360 6.366 31.508 36.186 13.729 35.74 MANMA1002887 53.445 53.672 210.646 36.086 31.508 36.186 13.729 35.74 MANMA1002883 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.	MANMA1002838	84.752						11.628	26.416
MAIMA1002843 76.343 31.051 107.479 18.190 24.282 30.455 19.401 13.72	MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.161 MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.161 MANMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MANMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.091 MANMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.98 MANMA1002868 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002875 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.10 MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 MANMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.41 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.14 MANMA1002887 45.505 7.809 7.546 6.360 6.366 31.508 36.186 13.729 35.74 MANMA1002887 53.445 53.672 210.646 36.086 31.508 36.186 13.729 35.74 MANMA1002883 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.		98.706	53.519	151.675	23. 902	32.033			47.227
MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.16 MANMA1002845 4.464 5.631 16.258 13.028 3.642 8.306 5.338 22.84 MANMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.86 MANMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.05 MANMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.91 MANMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 38.439 58.44 MANMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.14 MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.33 MANMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 MANMA1002887 45.505 7.809 7.548 7.024 9.965 8.271 13.675 5.154 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.154 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 38.649 41.0 MANMA1002887 53.475 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MANMA1002893 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2	MANMA1002844 311.853 139.150 228.560 66.881 72.282 201.758 152.946 94.161 MANMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.80 MANMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.09 MANMA1002868 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.98 MANMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 38.439 58.46 MANMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.10 MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 MANMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.41 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548						24. 282	30.456		13.727
MANIMA1002855	MANNA1002855							201,758		94. 166
MAIMA1002857 77.604 209.913 235.780 167.148 50.200 178.228 129.737 278.86 MAIMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.05 MAIMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.94 MAIMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 38.439 58.44 MAIMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.88 MAIMA1002867 28.097 6.998 5.660 1.623 3.087 7.477 5.467 3.44 MAIMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MAIMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MAIMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MAIMA1002879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.14 30.003	MANNA1002857									
MAIMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.09 MAIMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.90 MAIMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 38.439 58.40 MAIMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.83 MAIMA1002871 28.097 66.998 5.650 1.623 3.087 7.477 5.467 3.40 MAIMA1002872 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.90 MAIMA1002873 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.90 MAIMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.33 MAIMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 MAIMA1002885 87.039 28.425 35.323 14.016 29.952 14.101 61.975 26.2 MAIMA1002886 398.174 39.003 88.206 52.831 26.325 197.562 39.216 20.58 MAIMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 MAIMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 MAIMA1002887 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MAIMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2	MANMA1002858 113.809 319.730 662.654 523.500 84.144 532.413 382.518 1000.09 MANMA1002863 108.297 33.190 66.980 38.305 26.112 45.735 86.883 51.98 MANMA1002868 65.375 102.643 253.035 92.062 91.774 46.567 38.439 58.465 MANMA1002869 85.453 22.923 80.058 19.164 22.933 26.217 42.600 30.85 MANMA1002871 28.097 66.998 5.660 1.623 3.087 7.477 5.467 3.40 MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94 MANMA1002879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.10 MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 MANMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.41 MANMA1002885 87.039 28.425 35.323 14.016 29.952 34.101 61.975 26.27 MANMA1002886 398.174 39.003 88.206 52.831 26.325 197.562 39.216 20.566 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002887 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29									
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MANMATOO2871 28.097 6.998 5.660 1.623 3.087 7.477 5.467 3.4 MANMATOO2875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.9 MANMATOO2879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.16 MANMATOO2880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 MANMATOO2881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 MANMATOO2885 87.039 28.425 35.123 14.016 29.952 34.101 61.975 26.2 MANMATOO2886 398.174 39.003 88.206 52.831 26.325 197.562 39.216 20.5 MANMATOO2887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 MANMATOO2889 52.66 61.707 153.034 36.444 <	MANMA1002871 28.097 6.998 5.660 1.623 3.087 7.477 5.467 3.40	MAMMA1002869	85, 453	22, 923	80.058	19.164	22.933	26.217	42.600	30.859
MANMA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.96 MANMA1002879 33.352 14.773 9.446 6.359 8.506 13.275 30.077 23.16 MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 MANMA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 MANMA1002885 87.039 28.425 35.123 14.016 29.952 34.101 61.975 26.2 MANMA1002886 398.174 39.003 88.206 52.831 26.325 197.562 39.216 20.5 MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 MANMA1002890 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.0 MANMA1002892 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2	MANNA1002875 20.954 16.542 18.160 22.628 23.110 21.099 24.952 32.94									3.406
MANMA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952 14.016 29.952	MANNA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 14.848 22.560 23.33 27.378 18.964 34.053 52.41 18.848 23.847									32.949
HAMMAT002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.3 MAMMAT002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.4 MAMMAT002885 87.039 28.425 35.223 14.016 29.952 34.101 61.975 26.2 MAMMAT002886 398.174 39.003 88.206 52.831 26.325 197.562 39.216 20.5 MAMMAT002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 MAMMAT002890 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.0 MAMMAT002882 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MAMMAT002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2	MANNA1002880 46.288 35.830 71.009 12.119 12.813 15.447 20.107 22.35 MANNA1002881 57.225 55.154 238.977 25.333 27.378 18.964 34.053 52.41 MANNA1002885 87.039 28.425 35.323 14.016 29.952 34.101 61.975 26.27 MANNA1002886 398.174 39.003 88.206 52.831 26.325 197.562 39.216 20.566 MANNA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANNA1002890 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.02 MANNA1002892 58.445 53.672 210.646 36.086 31.508 36.186 13.729 35.74 MANNA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29									
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MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.1 MANMA1002880 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.0 MANMA1002892 58,445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2	MANMA1002887 45.505 7.809 7.548 7.024 9.968 8.271 13.675 5.11 MANMA1002890 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.02 MANMA1002892 58,445 53,672 210.646 36.086 31.508 36.186 13.729 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29			39.003		52.831		197.562	39, 216	20.561
MANMA1002890 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.0 MANMA1002892 58,445 53.672 210.646 36.086 31.508 36.186 13.729 35.7 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.2	MANNA1002890 65.426 61.707 153.034 36.444 19.739 40.974 38.649 41.02 MANNA1002892 58,445 53.672 210.646 36.086 31.508 36.186 13.729 35.74 MANNA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29									5. 111
MANMA1002892 58, 445 53, 672 210, 646 36, 086 31, 508 36, 186 13, 729 35, 7 MANMA1002893 76, 469 18, 593 25, 600 5, 864 9, 192 24, 826 20, 585 11, 2	MANMA1002892 58,445 53,672 210.646 36.086 31.508 36.186 13.729 35.74 MANMA1002893 76.469 18.593 25.600 5.864 9.192 24.826 20.585 11.29									
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Table 61

MANMA 1 DO	2898	88. 538	24. 524	42.725	9,653	16.551	32, 137	42. 359	30.615
MAMMA 100		191.445	39.095	72.714	28. 234	32.209	91.200	60.899	51.358
MANMA 100	2906	92.692	27.862	53.273	26.259	34. 130	57. 141	67.635	26. 917
MAMMA 1 CC	2908	77.656	66.964	209.054	54.014	54. 429	43.639	58. 626	50. 901
MANMA 1 OT	2909	157. 128	123.626		152.777	89. 304	83.884	61.550	89.879
MANNA 100		55. 362	26. 201	35. 298	14. 931	10.960	19. 166	27.775	29. 119
MAMMA 1 DO		50. 571	70.116	54. 395	18.071	27.814	43. 511	11.984	57. 467
MAMMA 100		105. 041	221.644	119.112	66.217	73.866		1218. 974	550. 265
MAMMA 100		68.089	38.713	147. 112	32. 243	19. 181	31.875	24. 698	46. 379
MAMMA 100		207. 866	61.711	89.764	38. 377	38.050 10.152	97.677 15.165	156.876 14.230	119. 279
MAMMA 100		34.139 18.884	13.727 30.845	21.350 50.805	7. 309 19. 591	7.699	16. 322	11. 528	24. 529
MAMMA 100		63.095	31.441	46.623	20. 590	18.624	28. 594	29. 987	39. 586
MANMA 10		43. 981	37.785	133.836	22, 173	11.661	25. 346	15. 389	28. 296
MANNA TO		37.974	16,689	23.126	13. 527	10.863	35.085	22.091	25.886
MAMMA 10		178.268	124. 368		120. 984	97.317	92.795	66.069	109.854
MAMMA 10	02971	99. 466	79.461	50.710	19.662	15.091	40.745	37. 592	51.546
MAMMA 10		83. 922	33.377	50.911	16. 436	12.354	42.113	50. 137	45. 819
MANMA 10		117. 540	70.913,	318.513	45. 601	38. 568	34.070	22.903	68.699
MAMMA 10		80.771	204.398	227. 280	56. 459	375.745	119.386	122.750	226. 538
MANMA 10		19.895	9. 493 50. 918	14. 202 156. 507	6. 265 28. 534	0.000 30.958	0.000 22.630	0.000 16.594	5.076 36.952
MAMMA10		65. 397 104. 891	69.630	125. 933	48.800	36, 915	48. 025	45.716	47. 346
MANNA 10		41.353	106.059		111.746	92.691	59. 597	33.719	77.654
MAMMA 10		20. 423	21.289	75.498	16.044	8.909	15.878	8. 947	15.193
MAMMA 10		45.615	37.641	29.754	23.843	21.157	33. 395	48. 907	39.054
MAMMA 10	03013	65.088	58. 284	49. 438	27. 289	18.877	31.768	67. 950	59.419
MANNA 10		36.817	29. 585	89. 251	19.826	4. 679	16, 502	6. 959	10, 432
MAJAMA 10		10.026	30.107	5. 244	7.467	2.375	6. 403	3. 225	6.184
MAJMA 10		48.046	31.761	50. 515	13. 842 8. 603	17. 142 6. 618	19.341 9.838	28.497 11.161	20. 218 6. 781
MAMMA 10 MAMMA 10		28.646 248.219	14. 274 140. 526	3.514 311.997	98. 494	105, 194	112.752	66. 462	132.570
MANMA 10		47.072	27. 208	130.132	44.811	42.096	33. 806	17, 555	36.757
MANMATO		102.528	49. 560	45.025	30. 912	25. 924	64. 046	42.175	56.246
NAMMA 10		37.382	19.822	98.219	37. 555	17. 115	27. 935	9, 656	25. 906
O I AMMAN	03040	76.014	95.416	243. 138	114.795	84.250	59. 989	42.107	100.448
MANMA 10		79, 444	46. 915	90.545	40. 709	21. 121	25. 258	13.745	23. 444
MAMMA 10		376.340	121.483	150.100	91.015	100. 397	168. 621	175. 219	122. 400
MANNA 10		26.899	9.631	9. 169	2. 907	5. 679 15. 758	12.149	5.016 6.277	10.003
MAMMA 10 MAMMA 10		38.639 31.238	24. 977 13. 811	76.695 32.121	21.811 15.345	7.891	17.689	3. 176	18.147
MAMMA 10		68. 258	35. 596	34.053	23.862	19.335	28. 373	32, 521	36.634
MAMMA 10		43.837	46.015	117.875	31.178	11.361	17.068	9. 179	35.831
MAMMA 10		16.366	6. 334	32.629	10.374	3.215	6. 507	2. 433	11.804
MAMMA 10		49.867	51.500	220.715	36. 189	24.057	14, 625	14, 530	41.852
MAMMA 10		22.129	73.102	15.615	27.304	11.693	9. 575	15. 986	84.963
MAMMA 1 C		8.240	37.313	24.078	8.354	10.123	9. 662	24. 609	12.392
MAMMA 1 C		44.094	27. 545 18. 730	96.117	16.060 14.500	12. 184 22. 389	15.519 16.929	4. 930 20. 089	23.720
MANIMATO MANIMATO		44. 491 35. 977	19, 146	31.447 34.647	14. 588	10, 720	11. 459	11.385	18. 999
MANMATO MANMATO		41.697	21.092	30, 337	15, 635	14.764	14, 690	17. 723	23.810
HAMMA 1		20.042	39. 595	102.916	21. 241	15. 167	17, 921	20.876	26.563
MAMMA 1		57.961	27.221	102.332	12.486	8.002	12. 295	13.773	22.285
MAMMA 1		267.516	37.924	129.263	66.561	86.667	135. 209	95. 293	83.256
MAMMA 1		22.855	14. 308	5.624	7. 938	2.690	14. 984	7.633	17. 269
MAMMA 1		6.575	9.140	33.040	4. 487	0.895	1.900	5.064	5.312
MANNA 1		14.105	18.018	18.562	11.213	11.461	16.500	8.591	9.815
MAMMA 1		311.806	87.992	58.938	77. 271	104.739	165. 139 31. 742	115.042 26.299	46.945 27.565
MAMMA 1		93.002	39.912 26.308	37.471 36.508	22.819 14.326	19.655	30.842	23.489	18.046
MANNA 1		41.709 34.876	32.317	147, 845	12, 108	24, 093	12.999	8.766	19.930
MAMMA		37.900	25. 338	29.052	18, 551	20.826	32.639	35. 893	33,749
MAMAT		26, 961	14.747	18.545		5. 852	14.778	13.694	20.137
MAMMAT	003166	12. 213	5. 478	7.671	8.749	1.781	3.094	8.412	7.640

Table 62

55

						T	15 100 T	16 400 I	11 550
	NB9N31000010	31.105	17.113	26. 284	14. 271	7. 540	17.180	16.220	11.568
	NB9N31000016	63.431	16, 195	24.879	17.001	16.740	25. 216	14.845	17.364
	NB9N31000043	87. 438	35, 161	58.144	20.813	36.473	36, 956	51.575	34.673
5									
	NB9N31000045	83. 399	109. 448	62.101	95.653	93.734	94.218	166.654	74.328
	NB9N31000054	41.821	12.636	37.831	15.025	15. 265	18.963	10.894	13. 189
	NB9N31000076	22.822	22, 709	57. 320	14. 223	12.517	9,029	11.713	24. 494
						11.744	29.951	13.909	30.012
	NB9N31000086	31.281	74.504	22.561	29. 164				
	NT2RM1000001	11.595	9. 900	11.540	4. 467	4.016	8.823	6.775	5.184
	NT2RM1000018	333.185	68.022	171, 103	77.680	48.418	138, 131	122.906	79.595
10			9. 768	23.088	9. 453	13. 222	16. 128	22.911	12.495
	NT2RM1000032	37.506							
	NT2RM1000035	185. 573	46.513	81.354	56.890	39. 846	82, 885	74. 450	52.553
	NT2RM1000037	185.843	60.878	116.479	50.830	36.658	98.591	49.882	54. 356
	NT2RM1000039	228.804	172.849	444.715	104, 606	82.108	214, 282	139.766	101.078
									184.618
	NT2RM1000042	55.479	102.774		145. 900	52.898	89.445	80.537	
	NT2RM1000055	1.083	0.593	0.000	0.000	0. 252	0.000	5. 227	0.000
15	NT2RM1000059	212.057	100.267	173.989	78. 130	50.792	143.445	83.189	102.504
					1. 925	2.705	2.434	25.015	10.555
	NT2RM1000062	11.755	9, 438	11.334					
	NT2RM1000065	153.505	42. 956	56.248	29.740	66.820	67.974	42.112	65. 531
	NT2RM1000066	26.794	6.539	7.914	2.716	6.609	8. 275	11.533	13.605
	NT2RM1000071			61.623	97.378	24.665	45.008	74. 491	266.252
		42.919	126.091						
	NT 2RM 1000080	12.803	1.714	1.023	4.022	2.135	8.919	13.254	4. 329
20	NT2RM1000086	393.857	146. 368	283.360	100.835	117.874	205. 973	155.085	102.325
	NT2RM1000092	12. 949	18.015	4. 187	6.602	2.600	0.000	5.579	17.636
				0,000	0.180	0.000	0.000	0,000	0.655
	NT2RM1000118	0.000	0. 276						
	NT2RM1000119	18.719	5. 828	9, 051	5. 794	3.873	5.048	19.700	10.812
	NT2RM1000121	2.231	0.000	7.566	3. 177	3.735	3. 309	1.697	3.614
	NT2RM1000122	309.647	84, 904	138. 129	58.379	75.966	213, 166	141.553	57.569
							4, 212	8. 594	5.786
25	NT2RM1000127	14. 133	3. 707	2. 380	2. 322	3.743			
	NT2RM1000131	1.661	1. 269	0.348	0.000	0.768	0.000	2. 271	2. 221
	NT2RM1000132	10.432	7.649	9, 599	3.479	7.287	11.592	13.046	10.752
	NT2RM1000153	39, 773	9, 302	10, 314	3.465	4.419	11.775	17, 131	12.503
							51.668		177.417
	NT2RM1000184	85. 966	171, 937	58.982	34. 486	22.674		129.969	
	NT2RM1000186	2.149	4.607	0.000	0.000	1.586	1. 226	3.974	7.121
	NT2RM1000187	29.354	12.303	16.019	17.222	15.020	17. 175	15, 232	18.703
30		16.274		17.316	6.834	4.725	5. 212	8.917	6.720
	NT2RM1000199		0.000						
	NT2RM1000213	17.361	14.639	43.481	9. 904	8.998	12. 127	6. 422	10.141
	NT2RM1000215	8.787	10,858	90.070	4.505	89.435	12.158	6.380	7.453
	NT2RM1000218	0.000	10.196	7, 239	2.227	1, 452	4. 273	8.324	4. 445
				0.000	47.537	20.172	44. 102	26.563	63.368
	NT2RM1000224	35.730	65. 418						
	NT2RM1000236	52.706	47. 803	20. 481	19. 138	42.513	21.813	58.118	100.492
35	NT2RM1000242	0,000	0,000	0.000	0.000	0.000	0.000	0.000	0.000
	NT2RM1000244	13.988	12.654	6, 957	9.937	6.047	8. 026	8, 938	3.968
					169.383	149. 200	192.609	174. 288	239.093
	NT2RM1000252	283.006	144, 305	358. 324					
	NT2RM1000256	284. 496	113.021	203. 771	67.954	94. 270	152.181	132. 435	150.452
	NT2RM1000257	8. 203	8.081	9,713	9.716	0.000	5.002	7.893	7.694
	NT2RM1000260	548. 461	312.072	494.663	164, 454	249.491	313.672	232.568	270.549
40	NT2RM1000269	9. 472	7.461	6.606	10.004	8.876	5.844	16.818	6.933
40									
	NT2RM1000271	8.917	1.259	3.857	2.440	2.317	4. 289	4. 982	5.727
	NT2RM1000272	83.425	97, 598	29. 246	80.462	22.650	25. 350	34, 266	157.515
	NT2RM1000273	27.031	19, 960	21,872	11. 127	5. 201	25. 896	29.976	17.270
			91, 340	28. 306	26. 224	11,534	34.723	32, 623	85, 440
	NT2RM1000274	42. 234							
	NT2RM1000280	14. 289	12. 359	21.912	7.205	7.361	10. 397	4. 200	10.119
45	NT2RM1000295	8. 249	4. 916	17.445	4.671	9.099	9. 454	2.185	[1.092]
40	MT2RM1000300	41.252	31, 172	52,474	15. 266	6.023	14.825	6.206	14. 221
			217.805	133.583	142, 504	77.271	155, 874	78.198	321.054
	NT2RM1000304	130.855							
	NT2RM1000314	255. 347	113. 392	165. 204	56.831	114.936	189. 937	108.461	113.313
	NT2RM1000318	4.002	22. 985	8.505	14. 343	0.836	6.124	14.391	25.194
	NT2RM1000335	10. 157	10.048	6,881	7.482	5.897	3.558	14.151	14.353
							10.884	5.578	6.704
50	NT2RM1000341	41.219	3.681	1.562	0.000	0.000			
55	NT2RM1000350	302.316	74. 071	106.873	34.040	61.895	149.078	112.517	85.201
			0.000	0,000	1.807	0.000	0.921	2.303	1.256
	NT2RM1000354	6.027	U. UUU						
	NT2RM1000354	6.027					249 162	14 695	225, 724
	NT2RM1000355	74. 352	158, 811	209.578	39, 101	103. 936	249.368	14.695	225.724
	NT2RM1000355 NT2RM1000361	74. 362 16. 299	158.811 10.575	209. 578 9. 446	39. 101 7. 432	103. 936 8. 424	7.383	4.356	5.053
	NT2RM1000355	74. 352	158.811 10.575	209. 578 9. 446	39, 101	103. 936 8. 424			

Table 63

	NYABULAAA224	04 504	0.016	10 007	00 701 1	22 22 1	61 670 1	** ***	21 704
	NT2RM1000372	93. 583	9.616	49.097	28.761	33.904	61.678	39, 147	31.524
	NT2RM1000377	42.186	17.871	22.783	12.668	13.142	15. 587	18. 377	23.602
5	NT2RM1000388	8. 811	19. 351	1. 155	5. 242	0.780	5. 795	6. 201	11.464
	NT2RM1000394	0.899	1.862	0.813	1.925	0. 438	0.000	0.000	0.000
	NT2RM1000399	1.641	5. 386	0.000	2. 270	0. 570	0.319	2.023	1. 257
	NT2RM1000407	69.180	19.536	39. 379	6.299	21.106	27. 229	14. 102	13.378
	NT2RM1000421	0.890	0.000	0.000	0.000	0. 456	0.150	0.000	0.000
	NT2RM1000422	102,028	152, 115		297.482	65. 137	134.344	50. 452	241.878
	NT2RM1000430	16.769	3. 286	12.402	4. 398	4. 506	12.149	11. 238	7.508
10	NT2RM1000462	167.815	117.695	165.008	62.828	65. 795	81.561	72.026	118.786
	NT2RM1000499	16.037	22.127	75. 152	12. 507	7.415	7. 335	41. 299	22.217
	NT2RM1000512	126.610	24. 122	12.786			46. 878	21, 802	
					25. 082	11.161			31.090
	NT2RM1000519	7.852	28.718	9. 178	14.716	6.756	27. 934	11.081	10.474
	NT2RM1000527	29.692	15. 338	24.471	17.418	45. 221	59. 291	31. 450	14.020
45	NT2RM1000539	14.790	19.300	31.135	14.824	2. 560	6.669	3.751	10.774
15	NT2RM1000542	118.560	38. 555	21.020	20.675	29. 849	30.176	22. 378	32.507
	NT2RM1000553	37. 329	18.841	47. 329	24. 533	23.901	33.590	34. 084	33.966
	NT2RM1000555	77.352	46.168	43.953	21.772	15.838	16.936	12.057	35.840
	NT2RM1000558	55.132	15. 424	20.508	7.987	7.249	8.886	23. 984	21.919
	NT2RM1000563	39.161	14.058	17.872	12.234	8.871	14. 324	12.341	13.462
	NT2RM1000566	3.172	7.323	0.000	2.755	1,243	3. 584	2.944	4.754
20	NT2RM1000570	65. 428	72.508	44. 124	24. 498	15. 164	26. 341	21.720	56.340
	NT2RM1000571	20.300	15.881	9.841	14. 197	7. 525	7. 964	16.668	9.893
	NT2RM1000574	45. 305	32.953	5.746	5.977	1.945	5.060	1.526	3.809
	NT2RM1000580	10.540	9. 295	12.139	8.734	2.114	6, 532	5. 687	7. 120
	NT2RM1000620	11,778	12.782	21.632	15. 504	5. 894	4. 488	3. 359	17. 303
	NT2RM1000623	3.914	2.515	0.416			0.715		
	NT2RM1000823				3. 125	0. 251		0.355	2.159
25	NT2RM1000630	17.633	6.091	6. 532	3.910	2.095	8. 257	7. 963	6.411
		5. 563	70.230	93.799	22.316	42.967	24.174	6.091	43.328
	NT2RN1000634	3. 427	3.869	2. 248	1.997	0.487	0.000	1.258	3.039
	NT2RM1000642	87.902	31.353	26.846	11.421	21.495	75. 074	66.152	42.393
	NT2RM1000647	45.410	65.742	56.619	55. 351	49. 439	30. 233	26.128	50.923
	NT2RM1000648	25. 285	9.969	8. 914	5. 538_	3. 383	6.086	5.045	5. 201
30	NT2RM1000650	22. 370	15.864	19.881	11.036	29.031	8. 360	13.836	11.166
••	NT2RM1000661	23. 325	6. 294	12.692	7. 551	6. 360	11.076	18.036	9.158
	NT2RM1000666	13.966	1.244	3. 221	1.629	1.543	4. 997	1.079	2.418
	NT2RM1000669	7.339	9.184	2.145	1.453	1.159	1.973	0.824	6.789
	NT2RM1000872	58.162	25.532	15. 778	9. 171	22.446	58. 987	16.791	14. 945
	NT2RM1000681	21.724	106.663	3.979	14.842	2.185	20. 284	16.034	21.688
	NT2RN1000691	4. 381	9. 202	2.832	3.483	1.268	0.878	2. 181	3.652
35	NT2RM1000698	31.943	17.379	9.609	16.495	5. 185	8.614	8. 628	12.092
	NT2RM1000699	10.439	2.722	5, 406	4.115	3. 535	6. 367	10.784	8.214
	NT2RM1000702	32.110	7.097	17.438	3.946	5.019	19.783	16.192	9.778
	NT2RM1000703	32.168	17.962	20.468	14.964	19.912	19.806	20. 940	16.286
	NT2RM1000704	25, 926	35.690	22.230	11.998	15.536	38.075	52.384	26.689
	NT2RM1000725	12.567	91,681	3.742	10.735	0. 262	10. 594	14.773	17.602
40	NT2RM1000726	7.525	9.354	5, 608	7.297	2. 528	3. 884	3. 237	8.489
	NT2RM1000731	144.609	19.850	46.338	14, 141	85.767	40. 231	32.791	30.972
	NT2RM1000741	14. 291	4.715	6. 122	2.576	3. 554	8. 230	5. 265	7.328
	NT2RM1000742	30.801	9. 241	6.240	6.116	3.655	11. 131	7.680	11.315
	NT2RM1000744	69.419	21.887	27.283	15.799	11, 433	38.093	24. 162	24.347
	NT2RM1000746	12.863	7.631	12.042	6.326	6.665	9. 321	8.974	11.118
	NT2RM1000747	24. 565	39, 958	11.215	5. 537	1.866	7.009	10.940	21.461
45	NT2RM1000752	13.148	7.585	3. 359	5.748	4. 905	1. 290	6.516	8.585
	NT2RM1000767	146.795	35. 621	33.719	11. 495	31.430	63. 425	41.576	22.788
	NT2RM1000770	24. 395			11. 954	11. 449	9, 412	14. 053	
			7.712	21.569					17.537
	NT2RM1000772	2.148	5. 100	1.271	2.181	0.000	1.505	6. 132	3.034
	NT2RM1000779	284. 561	185. 275	301.250	139.318	150. 250	196. 541	146. 279	96.926
50	NT2RM1000780	9. 227	9. 621	4. 260	6.864	3. 591	4. 298	8. 898	2.912
• •	NT2RM1000781	0.000	0.000	4.468	0.666	2. 562	3.064	2.407	2. 127
	NT2RM1000789	79.877	28. 387	74. 545	23.140	28. 956	35. 852	51.230	46. 548
	NT2RM1000800	4. 947	10.706	34.906	3.617	6.856	4. 436	8. 934	3. 531
	NT2RM1000802	209. 372	41.025	60.767	12.693	69.721	155. 310	133.291	27.049
	NT2RM1000811	0.000	0.807	0.000	3.615	0. 593	0.000	1.896	1.921

. Table 64

				. 16	1016 04				
	NT2RM1000826	55. 971	29.000	28.733	20.800	12. 255	7.195	28.144	23.708
	NT2RM1000829	39.377	19.978	34. 233	28.539	40.659	14.500	22. 956	26.065
	NT2RM1000831	92.244	176.233	212.504	115. 234	47. 485	121. 255		
5	NT2RM1000833	20.877	17.302	8.876	4. 821	8.474		114.428	264.692
	NT2RM1000834	7.920	13.742	7.973	9. 896		6.471	16. 424	13, 119
	NT2RM1000841	31,899				4.809	8.919	6.281	8. 562
	NT2RM1000848		32.922	28. 948	39.736	19.743	24.819	26.306	46.020
	NT2RM1000850	10.486	17. 213	11.047	9.143	7.207	4.310	8.632	18.858
		4.705	2.700	0.000	1.784	0.000	1. 597	2.104	7. 243
10	NT2RM1000852	27.699	10.440	14.655	3.679	11.796	13. 435	15.920	11.316
,,,	NT2RM1000853	0.000	4. 915	0.000	1.897	0.000	0.000	19.505	3.017
	NT2RM1000855	295.899	111.992	196.426	53. 443	65. 232	138. 673	132.776	97.678
	NT2RM1000857	419.515	279. 225	710.235	153. 528	198.222	264. 575	140.191	196.436
	NT2RM1000858	450. 537	223.032	628. 109	128. 574	92.997	272. 161	183.324	165.845
	NT2RM1000867	36.148	35. 491	71.518	26.137	22.828	37.610	46.674	48. 259
15	NT2RM1000874	94.766	25. 329	40.690	15. 917	33. 235	69.767	75.898	34.795
13	NT2RM1000882	32.751	18.077	43.528	12.957	13.381	12. 209	10.357	22.709
	NT2RM1000883	312.282	118. 317	233, 345	90. 226	109.110	311.111	130.746	182.823
	NT2RN1000885	252.089	146. 253	191, 597	129.087	63.370	152.039	156.686	193. 445
	NT2RM1000893	28.474	12.532,	13.539	21.087	13.367	23. 959	22.465	14.066
	NT2RM1000894	246.338	100. 240	188.863	51.822	48.537	189. 474	182.264	80.716
20	NT2RM1000898 NT2RM1000899	8.028	11.716	12.431	3. 461	8.055	10.349	3. 262	8.889
20	NT2RM1000905	20.978	2.796	3.034	4.018	6.936	7. 286	6.525	8.715
	NT2RM1000910	90.972	37.943	146. 214	36. 300	72.541	61.959	55. 239	46. 935
	NT2RM1000914	21.235	22.607	15. 176	6. 355	3.770	20. 204	15, 343	18.656
	NT2RM1000919	199. 944	90. 792	169. 446	46. 693	65. 449	122.556	87.145	72.117
	NT2RM1000919	36. 141	16. 161	19.116	13. 229	8.891	18.002	10.279	10. 389
05	NT2RM1000921	0.242	1.831	11.629	2.787	0.000	1.344	1.305	2. 292
25	NT2RM1000924	13.119 29.895	18.060 12.894	5. 555	12.140	3.037	3. 684	6.526	16.464
	NT2RM1000927	48.046	34.032	4. 946	4. 788	7. 984	10. 841	16.108	5. 749
	NT2RM1000951	13.349	11.379	49, 155 12, 531	23. 882	14.687	14. 867	17.603	20. 582
	NT2RM1000956	5. 337	16. 522	6. 739	13. 272 2. 246	6.919	7. 215	10.192	8. 882
	NT2RM1000960	24. 574	14. 841	49. 930	16.747	6.192	6. 379	6.215	8.675
20	NT2RM1000961	20. 594	16.610	28. 449	33.770	44. 584	52. 121	23.270	34.312
30	NT2RM1000962	1.479	8. 158	49. 309	6.863	11.295 4.421	30. 987 9. 226	65.017	30. 389
	NT2RM1000973	69. 241	51.561	16. 390	19. 560	15. 357	27.890	13.337	10. 246
	NT2RM1000978	0.000	0.000	0.000	0.000	0.000	0.000	33.675 0.000	45.410
•	NT2RM1000982	7. 275	2.308	2.120	2.059	1.138	1, 293	1.746	0.368
	NT2RM1000991	13.759	6.798	22, 345	7.467	6. 192	11, 494	1.942	4. 769 9. 039
35	NT2RM1000994	12.087	15, 119	14, 969	10.866	9. 132	2. 303	4, 549	14.654
35	NT2RM1001002	46. 263	5. 707	19, 271	15. 499	18.065	33. 283	21.225	33.831
•	NT2RM1001003	14. 107	33. 647	23.710	23.835	3. 391	10, 638	8. 307	14.681
	NT2RM1001008	4. 937	4. 696	0.740	4.466	2. 544	3, 192	3.215	10.971
	NT2RM1001011	67.834	16.031	21. 431	8.274	20. 203	46.979	40.030	18. 121
	NT2RM1001013	25. 323	6.694	3. 303	6.673	8.650	15, 882	23.168	23. 126
40	NT2RM1001017	8.644	4. 934	1.214	2, 455	1.873	2.894	4.062	7.068
70	NT2RM1001018	224.654	234,771	124.092	68,774	75.070	85.777	124.713	184.612
	NT2RM1001026	23. 853	12.510	10. 387	14, 301	5. 568	12.341	14.618	17.008
	NT2RM1001028	11.717	13, 271	17. 437	18.862	5.641	12.231	8. 930	11.443
	NT2RM1001043	21.614	13. 830	4. 261	8, 481	4.770	7.687	17.274	10.663
	NT2RN1001044	21.983	20. 272	44. 315	8, 181	4.171	5.809	4, 623	9. 566
45	NT2RM1001059	3. 169	2, 991	1.316	0.000	0.352	2.727	2.878	3.632
40	NT2RM1001063	0.879	5. 544	0.768	1,254	0.973	4. 181	1.761	5. 391
	NT2RM1001066	3.011	3, 061	0.000	3.241	0.000	1. 348	1. 228	3.011
	NT2RM1001072	13.706	7.601	5. 972	2.306	0.165	3. 139	5.672	5.851
	NT2RM1001074	32.455	14. 324	28. 723	10.090	6.573	10.841	7, 837	10. 538
	MT2RW1001076	7. 339	4. 891	0.792	2.511	0.000	5. 644	6.602	2.026
50	NT2RM1001082	63.705	50, 432	105. 417	34, 113	20. 331	17. 230	16. 378	21.799
	NT2RM1001085	13, 921	7. 236	4. 420	3.206	4. 563	0. 966	5. 984	4.704
	MT2RM1001092	16. 133	28.559	80. 293	36.442	13.840	23, 671	15, 948	30. 844
	NT 28M1001102	2. 299	0.000	0.000	0.000	0.000	2.006	1, 301	2.772
	NT 2RM 1001103	4, 293	14. 550	11.888	3.980	17.852	6. 345	2. 505	12. 387
	NT2RM1001105 NT2RM1001112	0,000	0.418	0.000	0.686	0.000	0.000	0.000	1.156
55	MI CAMIONITIS	6. 983	5, 403	12. 985	7.889	7.226	5. 412	8. 469	12.089

Table 65

				18	idle ob				
	NT2RM1001115	100.486	24.788	67. 251	18, 301	19, 421	53. 304	29.318	21.097
	NT2RM1001122	18.980	19, 515	19.938	11.109	10, 211	34. 308	33, 955	13. 422
	NT2RW1001136	4.811	3.751	2,520	1.126	0.765	2. 194	2.817	5, 117
5	NT2RM1001139	78.791	18, 931	27.710	8, 382	21.060	31.349	14.028	14.521
	NT2RM2000003	27.773	13. 438*-	12.296	3. 254	10. 288	4. 103	14. 697	22.880
	NT2RM2000006	64.154	36. 637	117.073	30. 277	27. 783	25. 842	17, 647	24. 349
	NT2RM2000010	57.806	33.217	60.148	20,749	86.788	23. 487	19.722	22. 551
	NT2RM2000013	24.877	27. 244	40.874	15. 590	40.045	30. 831	48. 932	35.344
	NT2RM2000030	68.595	26. 308	27.271	17. 595	26. 608	41. 165	43. 837	27. 939
10		22.984	13, 418	59.847		13.094	11.681	12. 137	11.426
	NT2RM2000032	35.892	5, 887		11.737 23.568	9. 740	51.053	23.006	23. 405
	NT2RM2000039			28. 101		5. 098	11, 101	20. 459	
	NT2RM2000042	7. 936	9, 200	20.886	10.060	5. 195	6.720	11, 105	10.744 5.712
	NT2RM2000092	12.085	11.085	15. 415	5.779				
	NT2RM2000093	51.998	31.271	57.365	24.041	26. 832	24. 640	12.930	20. 135
15	NT2RM2000101	34. 341	46.687	64. 294	27.692	29. 563	48. 487	33. 388	54. 246
	NT2RM2000104	73.163	48. 315	58. 786	33.739	39. 845	53.753	69, 151	73. 279
	NT2RM2000124	35.818	16. 923	31.954	10.723	11.012	23.770	21, 401	72. 254
	NT2RM2000155	31.139	23.019	27.033	12.467	9, 797	13.085	10. 315	17.050
	NT2RM2000191	151,075	54.651	87.171	59. 579	62.006	74. 514	126.950	91.326
	NT2RM2000192	0.760	2.690	0.971	4. 582	1. 137	2. 242	1.413	0.000
20	NT2RM2000239	92.578	36.060	71.933	31. 157	21. 570	60. 155	49.672	39. 127
20	NT2RM2000240	104.218	69. 966	77. 545	23. 453	53. 412	78, 029	64. 223	83, 906
	NT2RM2000241	70, 281	31, 167	42.733	18.007	14, 544	13, 456	26. 176	42. 298
	NT2RM2000250	72.366	22. 586	52. 512	23.631	19.076	29.100	50.616	50.848
	NT2RM2000259	90, 122	33.799	39. 931	17.198	9. 865	44.083	74. 558	29.086
	NT2RM2000260	340.036	40. 469	141.962	35.653	77.794	188.072	216.739	59. 426
	NT2RM2000265	24. 506	4. 177	38. 440	1. 951	3. 495	14. 217	14. 995	14. 683
25	NT2RM2000287	131.692	88.080	127. 535	51.611	38. 294	53. 574	55. 104	70. 583
	NT2RM2000306	45, 342	24, 950	44. 593	13.884	40. 471	40. 133	22.666	33. 254
	NT2RM2000312	13.383	57. 043	78.915	13. 258	60.055	90. 975	183.675	38. 391
	NT2RM2000322	33.318	18. 077	22. 354	11.030	6.002	8. 829	16. 962	15. 344
	NT2RM2000343	70.618	78, 514	302. 242	43. 179	64. 338	35. 838	84. 150	77. 161
••	NT2RM2000359	79, 203	25. 437	34, 945	19. 556	16. 348	47. 922	31.041	20. 663
30	NT2RW2000362	138.367	75.052	100.195	73.363	49. 276	128. 683	126.847	106. 528
	NT2RM2000363	41. 249	17.128	40. 363	12.316	18, 047	6. 982	11. 907	9. 239
	NT2RM2000368	225. 356	121, 451	100.718	49.727	89.663	128. 354	136.054	93. 203
	NT2RM2000371	88.897	208. 325	97. 848	212.525	33.081	80. 287	140.890	131.756
	NT2RM2000374	54. 398	55, 656	153.004	34. 316	25.750	36.072	34. 151	51. 955
	NT2RM2000387	31.537	35.012	44. 269	24. 245	23. 611	19.094	24. 288	26.745
35	NT2RM2000393	43.873	18.662	32.917	12.496	14. 167	17. 560	23. 452	33. 102
	NT2RM2000395	11.936	2. 901	3.145	1.722	4. 564	6, 102	4. 725	9. 257
	NT2RM2000402	26.540	28.616	42.681	18.209	10.970	24. 876	20.077	26. 993
	NT2RM2000405	29, 390	26. 302	56. 236	18. 391	18.624	17.673	19. 408	19. 435
	NT2RM2000407	213.973	77. 583	145. 459	42.798	73.678	124, 360	103. 989	122.635
	NT2RM2000410	46.375	23.782	29.096	10.711	13. 331	26.855	27. 992	20. 820
40	NT2RM2000420	41.781	29, 100	39.676	24.872	16.605	26.730	29. 136	43. 708
	NT2RM2000422	400, 274	145. 824	265.042	51.828	73.571	186.812	131, 563	125. 088
	NT2RM2000423	119,707	56. 563	272.757	58.213	50. 981	60. 353	42. 529	86. 903
	NT2RM2000452	44. 543	24. 735	36.727	13.780	10.160	32.134	23. 468	26.716
	NT2RM2000469	28.062	19.762	14.685	5.603	7.485		10.716	6. 249
	NT2RM2000490	57. 984	29. 556	42.743	16.403	19.316	36.503	21.105	31. 221
45	NT2RM2000497	44, 862	39. 966	107.651	23. 488	15. 277	19. 316	13.374	16, 412
	NT2RM2000502	49. 184	33.683	39.515	14.256	18.792	23. 598	23.921	27.778
	NT2RM2000504	53. 653	30. 376	46. 453	19.836	22. 257	39. 106	28. 508	19. 188
	NT2RM2000514	40.702	23. 938	23. 980	9.704	12.601	20. 319	19.147	27. 441
	NT2RM2000522	6.782	0.000	4.730	3.680	1.616	2.008	4, 021	14. 506
	NT 2RM2000540	28. 543	24. 938	24. 326	8.984	9. 799	16. 595	10.471	17. 045
50	NT2RM2000556	0.000	0.000	0.000	0.000	0.000	0.000	0.000	0.000
	NT2RM2000565	52.454	32, 231	48.697	17.373	14.758	42.730	24. 240	28. 218
	NT2RM2000566	31, 997	22. 486	34.598	11.793	7.665	32. 508	18, 105	35. 032
	NT2RM2000567	57.110	29, 153	45.058	10.738	15.606	44. 727	22. 394	28.766
	NT2RM2000569	113.652	91.632	187.867	40.645	36.420	58. 576	40. 151	50.117
	NT2RM2000577	61.308	16, 114	35. 195	12.694	14. 986	83.608	36. 221	60.695
55	NT2RM2000581	152.797	45, 271	66. 363	20.096	32.397	79. 582	62. 192	40.676
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Table 66

	NT2RM2000582	96.163	83.789	104.868	37.893	45.777	67.766	50. 428	50.631
	NT2RM2000588	109.847	89.480	119. 521	70.534	32. 168	143, 491	88. 984	95. 908
_	NT2RM2000589	91.130	45. 398	66. 143	21.774	22. 548	80.656	43.864	35. 379
5	NT2RM2000594	31.068	22.138	28.684	10.809	13. 325	34. 179	10.310	16.391
	NT2RM2000599	275. 423	132.063	221, 911	86.738	66. 363	237.294	209. 381	119.304
	NT2RM2000609	26.687	13.378	20.025	9.729	14. 321	19. 395	17. 956	8. 545
	NT2RM2000612	40.704	19.012	36. 338	9.471	15. 531	27.049	24.872	30. 269
	NT2RM2000622	45. 492	46. 307	46.012	27.097	17, 426	48. 495	30.090	42.927
	NT2RM2000623	279.041	219.374	245, 200	90.410	123.723	286. 194	221.925	144.950
10									
••	NT2RM2000624	52. 551	88.174	87.665	60. 273	35.044	29. 084	27.783	54.409
	NT2RM2000632	15. 461	13.673	11.853	13.378	8.044	7.114	6.910	5.808
	NT2RM2000635	24.726	21.442	42. 243	17.900	14. 353	23.119	10.306	20.675
	NT2RM2000836	45. 247	47.662	62.828	24. 460	33, 311	28.868	35.751	35. 343
	NT2RM2000639	34. 707	19.290	26. 594	15.919	12.875	28. 297	20. 526	11.317
	NT2RM2000649	39.662	37.102	62.088	31.152	32. 252	42. 335	77. 796 T	50.424
15	NT2RM2000658	53, 598	26.723	55. 360	19.176	26. 348	46.815	23.949	20.812
	NT2RM2000660	84. 441	62.193	66.364	13. 329	36.417	48. 267	23.694	40.215
	NT2RM2000669	17.352	23.877	38. 180	11.181	16.885	17. 594	13.008	20.479
•	NT2RM2000689	118.126	102.565,	102, 237	102.435	37.057	156. 147	96.539	140.413
	NT2RM2000691	29. 467	12.787	29.631	9.783	15. 294	28. 392	15.401	17. 161
00	NT2RM2000714	238.396	61.067	122. 264	38. 290	60.785	222. 914	188. 827	77.434
20	NT2RM2000718	9. 515	10.199	19.686	5.036	7. 922	8. 962	7. 572	22.010
	NT2RM2000732	44.022	24.869	42.915	12.209	29, 863	38. 537	30. 201	17.415
	NT2RM2000735	112.208	47.966	111, 282	57. 228	38. 980	78. 590	45, 888	59.237
							35. 449		
	NT2RM2000740	23. 990	62.438	143.286	24.030	26.159		22.001	29.845
	NT2RM2000743	15. 424	14.901	23.591	12.391	9.779	16. 339	8. 950	8.560
	NT2RM2000772	79.885	34, 020	54. 908	31.068	31.256	64.893	44. 735	55. 557
25	NT2RM2000773	56.846	36.465	77.155	26.645	32. 523	60.130	42.946	53.958
	NT2RM2000776	56.550	40.820	69.793	43.736	22. 285	89. 348	33. 285	45. 221
	NT2RM2000784	54.586	33.888	45. 181	19.559	21. 292	43. 103	25. 540	42.124
	NT2RM2000795	169.462	132.660	456, 283	117, 450	94.702	91, 566	59.832	91.914
	NT2RM2000796	12.942	12.033	20.129	5. 817	6.070	11.596	8. 538	11.009
	NT2RM2000798	67. 292	147. 984	71.980	42.802	43. 127	85. 427	63.126	132.706
30	NT2RM2000801	145. 709	152.451	160.956	85. 365	73.827	214. 221	157. 384	174.371
	NT2RM2000821	29.716	25. 994	36, 976	14. 293	9, 638	63.476	12. 133	3.427
	NT2RM2000829	77.695	36.834	148.015	32.077	69.569	70.012	26, 103	73.222
							48. 733	36.614	45.277
	NT2RM2000837	85.748	27.100	51.022	19. 432	22.405			
	NT2RM2000924	41.170	22.739	31.818_	6. 582	16.935	130, 595	55.870	42. 226
	MT2RM2000930	45. 514	31, 120	39.165	20.017	17. 433	49. 111	28. 135	30.171
35	NT2RM2000937	85. 092	19.912	28.613	13.728	34. 425	55. 176	53.959	15.755
				59.137	18. 909	23.056	57. 088	26. 370	29.465
	NT2RM2000939	63. 956	41.986						
	NT2RM2000942	141. 275	345.015	119.378	242.434	78. 282	274.472	112.054	436.171
	NT2RM2000951	32. 383	20.717	32.763	17.041	10.179	32.704	19.494	30.498
	NT2RM2000952	33, 160	18.882	34, 052	15. 194	27.783	44. 540	16.881	31.012
	NT2RM2000966	54.007	44, 546	57, 551	30. 397	27.965	78. 353	44. 947	77.916
				100.373			115.479	60.146	151.200
40	NT2RM2000973	96.188	97.082		31.654	38. 259			
	NT2RM2000983	66.024	27.357	40.970	16.277	25.768	44. 322	40. 901	34.882
	NT2RM2000984	38.635	39.635	42.628	14.734	10.729	39.002	24.661	39.000
	NT2RM2000994	38. 406	43, 907	36, 416	29. 496	24, 408	22. 384	18.679	31.517
	NT2RM2001004	74.509	45. 438	146.622	36.919	35.918	125. 242	81.529	92.360
	NT2RM2001022	195. 677	346.056	350, 501	243, 410	179. 341	419.711	214.981	540.668
45	NT2RM2001035	23, 201	26.826	34.867	15. 930	11.692	19.371	11.576	23.987
•	NT2RM2001038	18, 846	16.860	28.577	14. 251	9. 432	21. 182	12.726	12.544
				22,001	11.462	11.232	18. 219	25.898	31.106
	NT2RM2001043	31, 149	17. 293						
	NT2RM2001050	101.638	45.617	56.097	28. 126	32.674	61.600	49.621	79.938
	NT2RM2001055	83.075	29.856	49.927	15.739	32. 251	60, 461	35. 926	29.242
	NT2RM2001065	21.466	21.970	40.162	20.006	27.398	26. 370	15.034	14.433
							370. 161	257.064	228.430
50	NT2RM2001075	366. 658	258.334	337.690	128. 945	166. 931			
	MT2RM2001083	230, 683	79.913	107.950	30. 576	63.142	203. 365	79.590	24.253
	NT2RM2001100	182.772	114.627	137.289	65.878	54.062	141.899	155, 507	119.434
	NT2RM2001105	101.949	70.116	95, 624	50.863	39.812	104. 272	87.573	85.122
			<u> </u>						
	NT2RM2001109	48, 591	27. 328	30.825	11.569	12.495	53. 494	34. 958	45. 222
	NT2RM2001110	99. 871	68.967	152. 982	31.616	42.715	78.028	71.894	63.509
55	NT2RM2001126	57. 602	33.922	47.638	18.667	20.095	52.257	42.378	28. 204
	14. FUMEAA1 150	31.002	74. 35.	1	,	1 20.000		,	647

Table 67

	NT2RM2001131	59.454	21.547	32.934	24.063	22.706	37.676	28.873	17.418
	NT2RM2001141	116.250	82.599	275.090	51.756	53.614	85.069	47.274	63, 199
_	NT2RM2001152	20. 261	21.814	23. 297	10.506	9, 194	20.068	10.068	22.007
5	NT2RM2001177	44. 847	43. 449	52. 307	26.604	19. 552	41.709	26.283	55. 231
	NT2RM2001194	164.727	54. 905	97. 293	28. 358	44. 057	146.597	99.019	118.606
	NT2RM2001195	36.939	36. 245	34, 818	15, 750	15. 727	32.602	21.861	34. 274
	NT2RM2001196	125, 134	23. 362	52.729	15. 781	25.090	77.518	62.058	31.794
	NT2RM2001201	56. 981	42.504	62, 447	20. 139	31, 351	68.607	32.835	44, 422
40	NT2RM2001221	65.764	32.746	40. 357	19.556	25. 529	40.240	33, 849	36. 497
10	NT2RM2001238	34. 807	25. 200	33.023	13.254	14. 872	43.011	20, 155	18. 493
	NT2RM2001243	50. 316	49.076	42. 361	34 48	33. 121	68.021	35.734	60.810
	NT2RM2001244	39. 082	47.756	54.069	35. 242	30.728	59.908	22.778	50. 393
	NT2RM2001247	138. 825	184, 906	146. 564	65. 082	57.954	94. 133	78.544	136, 745
	NT2RM2001256	28. 147	18.773	29, 336	14. 133	9.881	8.739	16, 106	25. 473
45	NT2RM2001269	21.655	19.444	36, 676	14. 235	17.978	11.919	14. 441	17.847
15	NT2RM2001278	105. 133	67.683	225, 135	41.243	42.803	61.361	51.930	64. 103
	NT2RM2001291	21.264	19.798	31, 162	8.619	11.535	15.945	16. 243	12. 482
	NT2RM2001294	60.754	44. 696	66, 102	25. 820	20.715	42.950	28. 321	33. 134
	NT2RM2001295	43.856	35. 189 ,	40.675	10.220	16.301	35.694	20.908	35. 879
	NT2RM2001302	30.816	16.802	26.058	10.228	12. 245	25.513	14.404	12. 416
20	NT2RN2001306	11.584	52. 176	16.722	6.379	6.616	13.560	8.347	10. 145
20	NT2RM2001312	33. 361	18.866	54. 572	11.148	10.119	13.848	8. 526	26.714
	NT2RM2001319	13. 127	22. 841	- 23, 586	17.119	10. 492	18. 998	4. 495	36. 587
	NT2RM2001324	103. 673	83.091	165, 198	32.861	22.836	56.112	31.793	39. 459
	NT2RM2001345	49. 534	25.168	35, 284	14.837	16. 900	100.618	25. 540	19.919
	NT2RM2001360	74. 152	33.097	38. 122	17.360	16.021	50. 562	31. 265	21.915
25	NT2RM2001370	28.821	12.859	21.986	6. 327	5. 734	26.406	10.631	2. 394
20	NT2RM2001391	16. 127	5.412	27.834	4. 575	4, 553	14. 188	3.910	9. 994
	NT2RM2001393	57. 930	25. 241	58.135	14. 781	20. 544	47.187	32.903	28. 104
	NT2RM2001420	17. 272	10.676	16.079	6.774	6.751	2.717	3.157	8. 464
	NT2RM2001423	17. 345	9.837	15. 261	12. 233	6. 527	15. 432	10.007 95.111	10. 935
	NT2RM2001424 NT2RM2001482	196. 973	74.966 123.493	136.019 274.926	35. 222 59. 811	48. 814 62. 022	142. 268 227. 572	99.155	56. 187 72. 372
30	NT2RM2001402	65. 942	48.790	62. 383	28.605	19.730	68. 321	23.722	26. 475
	NT2RM2001504	39. 282	24.742	30, 958	9. 395	16.991	46.880	13.034	16, 709
	NT2RM2001524	24.755	14, 244	24. 384	9.699	10.204	16. 924	9.647	14. 539
	NT2RM2001530	5. 573	8.914	10.768	5.856	3. 286	9.623	4, 337	7. 511
	NT2RM2001533	69. 137	57.026	127.055	29.970	34.159	33.371	27.483	25. 268
	NT2RN2001540	65. 400	54. 541	73.017	63.277	35. 636	49.097	31.308	76.346
35	NT2RM2001544	18.067	19.624	25, 228	12.549	7.049	19.380	11.033	9. 485
	NT2RM2001547	22.357	25.608	19.122	11.755	13.130	14.503	12.339	10.697
	NT2RM2001558	59. 623	25.861	31.696	14, 111	16. 568	53.758	34, 606	18. 325
	NT2RM2001575	53. 128	46.425	111.368	27. 392	24. 257	43.005	25. 405	24. 423
	NT2RM2001582	59.050	42.778	132, 294	24. 555	24. 449	28.347	22. 303	22. 397
	NT2RM2001588	35. 342	21.815	27. 343	8.806	14. 132	21.498	16.451	22.464
40	NT2RM2001592	19. 456	18. 542	28. 436	10. 182	12. 538	15. 234	15.478	15. 460
	NT2RM2001603	42.456	15. 253	41.037	12.377	16.738	23.117	21.517	12. 277
	NT2RM2001605	60. 434	36.233	43. 204	13.580	20.116	41. 260	19.191	15. 459 16. 136
	NT2RM2001611 NT2RM2001613	54.771 39.500	39.056 22.894	128. 984 27. 579	17. 180	11.577	26.696	21.149	24.773
	NT2RM2001626	202.358	40.774	93. 458	19.731	45. 138	168. 993	96.729	42.842
45 '	NT2RM2001632	30. 150	45. 268	47. 586	25, 780	18.848	32.974	21.939	45, 513
40	NT2RM2001633	6. 521	9. 885	12.546	7.571	6.017	11. 226	7. 294	20. 798
	NT2RM2001635	188. 515	41.783	101. 462	30.227	41.863	115.049	88. 246	58.313
	NT 2RM2001636	26. 880	23.087	31.788	15.679	14. 225	22. 589	16.870	26. 264
	NT2RM2001637	13.020	5. 524	6.631	4.897	11.170	10.700	20.526	5. 331
	NT2RM2001639	71.531	28.740	32.389	12.149	15.813	54.897	28.931	13.443
50	NT2RM2001641	39. 297	32.462	49. 334	14.630	22.002	30.556	21.763	16.776
	NT2RM2001643	25. 535	12.621	15.764	6.658	12.027	21.274	22.136	12.847
	NT2RM2001648	26. 584	18.351	24. 507		6.636	18.218	14.277	13. 561
	NT2RM2001652	18.655	15.854	22. 304		9.644	25.729	7.851	20.144
	NT2RM2001659	16.893	10.861	16.538	3.750	4. 964	9. 228	6.172	11.278
	NT 2RM2001660	17.414	13. 987	20.619		10.544	12. 482	10.671	11. 244
55	NT2RM2001664	32.470	29. 186	27.804	16. 171	15. 728	29, 928	13.136	17.877

Table 68

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I.N	T2RM2001668	89.325	61.356	52. 221	32.055	34. 144	88.196	46.704	35. 968
TX	T2RM2001670	58.448	20. 552	40. 552	17.717	15. 452	67.725	25. 514	50. 962
	T2RM2001671	31, 368	15.752	21.018	19.630	8.980	62.746	15. 913	35. 807
	T2RM2001675	7. 281	7.210	6.726	2.026	5.059	4. 678	4.675	6.219
The state of the s	T2RM2001581	6.784	7.472	11.234	3. 584	7.454	5.095	5. 298	21. 547
îî	NT2RM2001685	28.752	21.105	22. 146	9. 525	9.058	19. 334	21. 485	26.746
	NT2RM2001688		25, 279	43.734	11. 154	11.656	30. 491	20. 238	33. 991
		35.233							
	NT2RM2001695	82.068	103.403	239. 543	46.305	60. 347	69. 201	35. 903	61.706
ħ	NT2RM2001696	101.355	65.027	68. 822	31.652	35, 701	110.799	51.358	52. 359
	NT2RM2001698	146.791	45. 033	83.099	28. 886	31.134	111.891	66.042	79. 232
	NT2RM2001699	24.737	20.994	25. 919	13.654	8. 984	19.132	14. 447	42. 307
- 1	NT2RM2001700	14.734	8.383	12. 975	3.702	1.312	7.813	9. 485	4. 374
Ī	NT2RM2001704	50.393	27.867	50.059	14, 943	24. 129	36.190	27.006	62. 522
	NT2RM2001706	75. 476	62,308	144.702	48. 167	41.996	55, 679	28. 916	57, 741
			12.916	25. 654	7. 345	10.946	13. 341	5.670	10. 957
	NT2RM2001714	14.876							
Į.	NT2RM2001716	294.058	99.615	122.970	48. 569	68.313	188. 154	109. 177	, 48. 112
П	NT2RM2001718	109.052	48.161	57.895	15, 717	34.379	105. 548	62.864	26.050
ħ	NT2RM2001723	20. 352	14. 923	16.575	7.233	8. 937	39. 809	8.807	9. 491
	NT2RM2001727	57.044	41.045	46. 272	22.665	16.545	51.332	33, 590	46. 539
	NT2RM2001730	27.206	22.859	24.865	8. 552	9. 397	19. 553	13.897	12. 427
	NT2RM2001738	25. 036	6. 229	18.054	7.967	10. 452	22. 532	14. 238	26.610
ħ	NT2RM2001743	31,219	15. 575	27.495	8.999	13.856	19.966	21. 123	40. 203
	NY2RM2001753	41.699	57.379	66.833	29. 155	36.474	48.608	37.342	50. 583
		102.308		95. 880	48. 800	50.926	85.016	46. 946	58. 535
	NT2RM2001755		95. 543						
L	NT2RM2001760	36.852	29. 592	43. 280	11.529	16. 235	41.973	21.095	36.897
	NT2RM2001765	17.310	22. 525	20.809	5. 472	6. 161	36. 420	11.083	21. 129
Ī	NT2RM2001767	507.383	198.624	244.752	82. 225	86.662	313.630	261.579	156. 449
	NT2RM2001768	14. 334	16.852	22. 405	14.516	7.327	13.653	4, 371	27.736
	NT2RM2001771	33.884	31.815	59.888	20. 959	19. 261	40.662	26. 114	70. 587
	NT2RM2001778	14.653	9.177	12.741	0.999	6.577	9. 552	8.651	6. 525
	NT2RM2001782	49. 540	17.667	39. 944	11.809	19.235	60. 433	38. 302	42.078
i	NT2RM2001784	31.529	23.807	34. 905	9.620	16.512	26.774	14.749	17.008
ì	NT2RM2001785	73.444	32.799	54.722	14, 868	28.332	74. 431	52.678	40. 155
	NT2RM2001792	82.550	48.689	54.661	13.880	26.470	67.309	56. 934	51, 170
	NT2RM2001795	130.534	65. 803	79. 887	22. 935	40.781	108. 971	66.672	68. 900
	NT2RM2001797	17.770	23. 911	46.302	31.918	15.965	38. 330	15. 267	61.440
	NT2RM2001800	32.076	15.750	32.039	9. 323	10.196	25. 569	24. 848	32. 579
1	NT2RM2001803	18.883	19.806	27.862	15. 915	15.790	17.317	12. 178	25.827
	NT2RM2001805	10.973	6.105	12.362	3.395	7.748	17.242	7.464	10.576
						14. 554	35. 269	18. 192	22. 416
	NT2RM2001806	41.604	28.683	30.345	12.360				
	NT2RM2001813	11.155	10. 752	12. 187	5. 926	6. 671	17.463	7.004	10.764
j	NT2RM2001814	15.422	18. 276	19.059	5. 168	10.179	14. 993	12. 571	9. 506
	NT2RM2001818	37.340	15.047	25. 378	7.050	13.614	28.082	23. 903	16.747
	NT2RM2001823	13.814	13. 268	12.712	4. 562	7.791	10.847	8.727	7.819
	NT2RM2001825	27.524	37. 936	22. 505	15. 145	17. 486	21.050	17. 161	33. 945
1	MIENMEUU 1843								
	NT2RM2001832	68.657	29.677	30. 202	9.749	22.522	37. 241	30.727	18. 205
	NT2RM2001839	53.715	31.908	39. 273	13. 944	12.144	27. 291	25. 952	18.816
	NT2RM2001840	108.411	98, 429	259.021	48.048	32.857	58.314	28. 523	37. 338
	NT2RM2001851	52, 202	39.752	63.088	24, 308	18.778	32.821	26.626	85.666
	NT2RM2001855	33.026	24, 176	29. 953	16. 912	19.394	23.562	31, 355	25.910
						16.183	32.799	17. 562	46.800
	NT2RM2001867	30, 838	22.957	35. 457	14. 948				
	NT2RM2001869	129. 599	162.083		173.694	64.737	231.277	145. 176	147. 129
	NT2RM2001879	14.477	14.016	20.104	6. 241	7.997	18. 463	6, 634	17. 934
	NT2RM2001883	42.649	14. 914	59.041	11.657	28.809	14.670	17. 172	5.396
	NT2RM2001886	31.621	19.917	31.650	19.861	14.683	19. 396	24.519	18.912
			18. 787	31.384	9, 308	6, 192	7.945	11.032	6.537
	NT2RM2001887	19, 995							
	MT2RM2001896							5347.627	1306. 593
	NT2RM2001902	9. 512	5176	9.030	3. 230	3. 539	7.418	7. 583	3. 383
	NT2RM2001903	63. 243	40.127	55. 162	28.793	22.732	77.356	28, 595	48. 438
	NT2RM2001930		64. 649	109. 195	31, 339	39.123	80.005	62, 289	59. 426
		108. 255							10.042
	NT2RM2001935	36.519	23. 148	18.415	4. 134	11.165	15. 562	19. 141	
	NT2RM2001936	78.536	47. 939	51.879	18. 980	21.013	48. 576	35. 554	52.419
	NT2RM2001939	23.961	5. 651	21, 192	6.301	5.377	17.197	7.025	5. 482
	NT2RM2001941	71.450	49.630	78. 923	19.738	22.274	54. 128	31.260	34.949
	1 PE 00 TENT 10	11.430	73.030	1	1 10.100	1 - 55 - 61 7	1 411.50		

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